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(54) Title: COMPOSITIONS AND METHODS FOR THERAPY AND DIAGNOSIS OF PROSTATE CANCER

## (57) Abstract

Compositions and methods for the therapy and diagnosis of cancer, such as prostate cancer, are disclosed. Compositions may comprise one or more prostate tumor proteins, immunogenic portions thereof, or polynucleotides that encode such portions. Alternatively, a therapeutic composition may comprise an antigen presenting cell that expresses a prostate tumor protein, or a T cell that is specific for cells expressing such a protein. Such compositions may be used, for example, for the prevention and treatment of diseases such as prostate cancer. Diagnostic methods based on detecting a prostate tumor protein, or mRNA encoding such a protein, in a sample are also provided.

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## COMPOSITIONS AND METHODS FOR THERAPY AND DIAGNOSIS OF PROSTATE CANCER

### TECHNICAL FIELD

The present invention relates generally to therapy and diagnosis of cancer, such as prostate cancer. The invention is more specifically related to polypeptides comprising at least a portion of a prostate tumor protein, and to polynucleotides encoding such polypeptides. Such polypeptides and polynucleotides may be used in vaccines and pharmaceutical compositions for prevention and treatment of prostate cancer, and for the diagnosis and monitoring of such cancers.

### BACKGROUND OF THE INVENTION

Prostate cancer is the most common form of cancer among males, with an estimated incidence of 30% in men over the age of 50. Overwhelming clinical evidence shows that human prostate cancer has the propensity to metastasize to bone, and the disease appears to progress inevitably from androgen dependent to androgen refractory status, leading to increased patient mortality. This prevalent disease is currently the second leading cause of cancer death among men in the U.S.

In spite of considerable research into therapies for the disease, prostate cancer remains difficult to treat. Commonly, treatment is based on surgery and/or radiation therapy, but these methods are ineffective in a significant percentage of cases. Two previously identified prostate specific proteins - prostate specific antigen (PSA) and prostatic acid phosphatase (PAP) - have limited therapeutic and diagnostic potential. For example, PSA levels do not always correlate well with the presence of prostate cancer, being positive in a percentage of non-prostate cancer cases, including benign prostatic hyperplasia (BPH). Furthermore, PSA measurements correlate with prostate volume, and do not indicate the level of metastasis.

In spite of considerable research into therapies for these and other cancers, prostate cancer remains difficult to diagnose and treat effectively. Accordingly, there is a need in the art for improved methods for detecting and treating such cancers. The present invention fulfills these needs and further provides other related advantages.

### SUMMARY OF THE INVENTION

Briefly stated, the present invention provides compositions and methods for the diagnosis and therapy of cancer, such as prostate cancer. In one aspect, the present

invention provides polypeptides comprising at least a portion of a prostate tumor protein, or a variant thereof. Certain portions and other variants are immunogenic, such that the ability of the variant to react with antigen-specific antisera is not substantially diminished. Within certain embodiments, the polypeptide comprises at least an immunogenic portion of a prostate tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of: (a) sequences recited in any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472; (b) sequences that hybridize to any of the foregoing sequences under moderately stringent conditions; and (c) complements of any of the sequence of (a) or (b). In certain specific embodiments, such a polypeptide comprises at least a portion, or variant thereof, of a tumor protein that includes an amino acid sequence selected from the group consisting of sequences recited in any one of SEQ ID NO: 112-114, 172, 176, 178, 327, 329, 331, 336, 339, 376-380 and 383.

The present invention further provides polynucleotides that encode a polypeptide as described above, or a portion thereof (such as a portion encoding at least 15 amino acid residues of a prostate tumor protein), expression vectors comprising such polynucleotides and host cells transformed or transfected with such expression vectors.

Within other aspects, the present invention provides pharmaceutical compositions comprising a polypeptide or polynucleotide as described above and a physiologically acceptable carrier.

Within a related aspect of the present invention, vaccines are provided. Such vaccines comprise a polypeptide or polynucleotide as described above and a non-specific immune response enhancer.

The present invention further provides pharmaceutical compositions that comprise: (a) an antibody or antigen-binding fragment thereof that specifically binds to a prostate tumor protein; and (b) a physiologically acceptable carrier.

Within further aspects, the present invention provides pharmaceutical compositions comprising: (a) an antigen presenting cell that expresses a polypeptide as described above and (b) a pharmaceutically acceptable carrier or excipient. Antigen presenting cells include dendritic cells, macrophages, monocytes, fibroblasts and B cells.

Within related aspects, vaccines are provided that comprise: (a) an antigen presenting cell that expresses a polypeptide as described above and (b) a non-specific immune response enhancer.

The present invention further provides, in other aspects, fusion proteins that comprise at least one polypeptide as described above, as well as polynucleotides encoding such fusion proteins.

Within related aspects, pharmaceutical compositions comprising a fusion protein, or a polynucleotide encoding a fusion protein, in combination with a physiologically acceptable carrier are provided.

Vaccines are further provided, within other aspects, that comprise a fusion protein, or a polynucleotide encoding a fusion protein, in combination with a non-specific immune response enhancer.

Within further aspects, the present invention provides methods for inhibiting the development of a cancer in a patient, comprising administering to a patient a pharmaceutical composition or vaccine as recited above.

The present invention further provides, within other aspects, methods for removing tumor cells from a biological sample, comprising contacting a biological sample with T cells that specifically react with a prostate tumor protein, wherein the step of contacting is performed under conditions and for a time sufficient to permit the removal of cells expressing the protein from the sample.

Within related aspects, methods are provided for inhibiting the development of a cancer in a patient, comprising administering to a patient a biological sample treated as described above.

Methods are further provided, within other aspects, for stimulating and/or expanding T cells specific for a prostate tumor protein, comprising contacting T cells with one or more of: (i) a polypeptide as described above; (ii) a polynucleotide encoding such a polypeptide; and/or (iii) an antigen presenting cell that expresses such a polypeptide; under conditions and for a time sufficient to permit the stimulation and/or expansion of T cells. Isolated T cell populations comprising T cells prepared as described above are also provided.

Within further aspects, the present invention provides methods for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a T cell population as described above.

The present invention further provides methods for inhibiting the development of a cancer in a patient, comprising the steps of: (a) incubating CD4<sup>+</sup> and/or CD8<sup>+</sup> T cells isolated from a patient with one or more of: (i) a polypeptide comprising at least an immunogenic portion of a prostate tumor protein; (ii) a polynucleotide encoding such a polypeptide; and (iii) an antigen-presenting cell that expressed such a polypeptide; and (b) administering to the patient an effective amount of the proliferated T cells, and thereby inhibiting the development of a cancer in the patient. Proliferated cells may, but need not, be cloned prior to administration to the patient.

Within further aspects, the present invention provides methods for determining the presence or absence of a cancer in a patient, comprising: (a) contacting a biological sample obtained from a patient with a binding agent that binds to a polypeptide as recited

above; (b) detecting in the sample an amount of polypeptide that binds to the binding agent; and (c) comparing the amount of polypeptide with a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient. Within preferred embodiments, the binding agent is an antibody, more preferably a monoclonal antibody. The cancer may be prostate cancer.

The present invention also provides, within other aspects, methods for monitoring the progression of a cancer in a patient. Such methods comprise the steps of: (a) contacting a biological sample obtained from a patient at a first point in time with a binding agent that binds to a polypeptide as recited above; (b) detecting in the sample an amount of polypeptide that binds to the binding agent; (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and (d) comparing the amount of polypeptide detected in step (c) with the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

The present invention further provides, within other aspects, methods for determining the presence or absence of a cancer in a patient, comprising the steps of: (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a prostate tumor protein; (b) detecting in the sample a level of a polynucleotide, preferably mRNA, that hybridizes to the oligonucleotide; and (c) comparing the level of polynucleotide that hybridizes to the oligonucleotide with a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient. Within certain embodiments, the amount of mRNA is detected via polymerase chain reaction using, for example, at least one oligonucleotide primer that hybridizes to a polynucleotide encoding a polypeptide as recited above, or a complement of such a polynucleotide. Within other embodiments, the amount of mRNA is detected using a hybridization technique, employing an oligonucleotide probe that hybridizes to a polynucleotide that encodes a polypeptide as recited above, or a complement of such a polynucleotide.

In related aspects, methods are provided for monitoring the progression of a cancer in a patient, comprising the steps of: (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a prostate tumor protein; (b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide; (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and (d) comparing the amount of polynucleotide detected in step (c) with the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

Within further aspects, the present invention provides antibodies, such as monoclonal antibodies, that bind to a polypeptide as described above, as well as diagnostic

kits comprising such antibodies. Diagnostic kits comprising one or more oligonucleotide probes or primers as described above are also provided.

These and other aspects of the present invention will become apparent upon reference to the following detailed description and attached drawings. All references disclosed herein are hereby incorporated by reference in their entirety as if each was incorporated individually.

## BRIEF DESCRIPTION OF THE DRAWINGS AND SEQUENCE IDENTIFIERS

Figure 1 illustrates the ability of T cells to kill fibroblasts expressing the representative prostate tumor polypeptide P502S, as compared to control fibroblasts. The percentage lysis is shown as a series of effector:target ratios, as indicated.

Figures 2A and 2B illustrate the ability of T cells to recognize cells expressing the representative prostate tumor polypeptide P502S. In each case, the number of  $\gamma$ -interferon spots is shown for different numbers of responders. In Figure 2A, data is presented for fibroblasts pulsed with the P2S-12 peptide, as compared to fibroblasts pulsed with a control E75 peptide. In Figure 2B, data is presented for fibroblasts expressing P502S, as compared to fibroblasts expressing HER-2/*neu*.

Figure 3 represents a peptide competition binding assay showing that the P1S#10 peptide, derived from P501S, binds HLA-A2. Peptide P1S#10 inhibits HLA-A2 restricted presentation of fluM58 peptide to CTL clone D150M58 in TNF release bioassay. D150M58 CTL is specific for the HLA-A2 binding influenza matrix peptide fluM58.

Figure 4 illustrates the ability of T cell lines generated from P1S#10 immunized mice to specifically lyse P1S#10-pulsed Jurkat A2Kb targets and P501S-transduced Jurkat A2Kb targets, as compared to EGFP-transduced Jurkat A2Kb. The percent lysis is shown as a series of effector to target ratios, as indicated.

Figure 5 illustrates the ability of a T cell clone to recognize and specifically lyse Jurkat A2Kb cells expressing the representative prostate tumor polypeptide P501S, thereby demonstrating that the P1S#10 peptide may be a naturally processed epitope of the P501S polypeptide.

Figures 6A and 6B are graphs illustrating the specificity of a CD8<sup>+</sup> cell line (3A-1) for a representative prostate tumor antigen (P501S). Figure 6A shows the results of a <sup>51</sup>Cr release assay. The percent specific lysis is shown as a series of effector:target ratios, as indicated. Figure 6B shows the production of interferon-gamma by 3A-1 cells stimulated with autologous B-LCL transduced with P501S, at varying effector:target ratios as indicated.

SEQ ID NO: 1 is the determined cDNA sequence for F1-13

SEQ ID NO: 2 is the determined 3' cDNA sequence for F1-12

SEQ ID NO: 3 is the determined 5' cDNA sequence for F1-12  
SEQ ID NO: 4 is the determined 3' cDNA sequence for F1-16  
SEQ ID NO: 5 is the determined 3' cDNA sequence for H1-1  
SEQ ID NO: 6 is the determined 3' cDNA sequence for H1-9  
SEQ ID NO: 7 is the determined 3' cDNA sequence for H1-4  
SEQ ID NO: 8 is the determined 3' cDNA sequence for J1-17  
SEQ ID NO: 9 is the determined 5' cDNA sequence for J1-17  
SEQ ID NO: 10 is the determined 3' cDNA sequence for L1-12  
SEQ ID NO: 11 is the determined 5' cDNA sequence for L1-12  
SEQ ID NO: 12 is the determined 3' cDNA sequence for N1-1862  
SEQ ID NO: 13 is the determined 5' cDNA sequence for N1-1862  
SEQ ID NO: 14 is the determined 3' cDNA sequence for J1-13  
SEQ ID NO: 15 is the determined 5' cDNA sequence for J1-13  
SEQ ID NO: 16 is the determined 3' cDNA sequence for J1-19  
SEQ ID NO: 17 is the determined 5' cDNA sequence for J1-19  
SEQ ID NO: 18 is the determined 3' cDNA sequence for J1-25  
SEQ ID NO: 19 is the determined 5' cDNA sequence for J1-25  
SEQ ID NO: 20 is the determined 5' cDNA sequence for J1-24  
SEQ ID NO: 21 is the determined 3' cDNA sequence for J1-24  
SEQ ID NO: 22 is the determined 5' cDNA sequence for K1-58  
SEQ ID NO: 23 is the determined 3' cDNA sequence for K1-58  
SEQ ID NO: 24 is the determined 5' cDNA sequence for K1-63  
SEQ ID NO: 25 is the determined 3' cDNA sequence for K1-63  
SEQ ID NO: 26 is the determined 5' cDNA sequence for L1-4  
SEQ ID NO: 27 is the determined 3' cDNA sequence for L1-4  
SEQ ID NO: 28 is the determined 5' cDNA sequence for L1-14  
SEQ ID NO: 29 is the determined 3' cDNA sequence for L1-14  
SEQ ID NO: 30 is the determined 3' cDNA sequence for J1-12  
SEQ ID NO: 31 is the determined 3' cDNA sequence for J1-16  
SEQ ID NO: 32 is the determined 3' cDNA sequence for J1-21  
SEQ ID NO: 33 is the determined 3' cDNA sequence for K1-48  
SEQ ID NO: 34 is the determined 3' cDNA sequence for K1-55  
SEQ ID NO: 35 is the determined 3' cDNA sequence for L1-2  
SEQ ID NO: 36 is the determined 3' cDNA sequence for L1-6  
SEQ ID NO: 37 is the determined 3' cDNA sequence for N1-1858  
SEQ ID NO: 38 is the determined 3' cDNA sequence for N1-1860  
SEQ ID NO: 39 is the determined 3' cDNA sequence for N1-1861

SEQ ID NO: 40 is the determined 3' cDNA sequence for N1-1864  
SEQ ID NO: 41 is the determined cDNA sequence for P5  
SEQ ID NO: 42 is the determined cDNA sequence for P8  
SEQ ID NO: 43 is the determined cDNA sequence for P9  
SEQ ID NO: 44 is the determined cDNA sequence for P18  
SEQ ID NO: 45 is the determined cDNA sequence for P20  
SEQ ID NO: 46 is the determined cDNA sequence for P29  
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SEQ ID NO: 64 is the determined cDNA sequence for P79  
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SEQ ID NO: 66 is the determined cDNA sequence for P68  
SEQ ID NO: 67 is the determined cDNA sequence for P80  
SEQ ID NO: 68 is the determined cDNA sequence for P82  
SEQ ID NO: 69 is the determined cDNA sequence for U1-3064  
SEQ ID NO: 70 is the determined cDNA sequence for U1-3065  
SEQ ID NO: 71 is the determined cDNA sequence for V1-3692  
SEQ ID NO: 72 is the determined cDNA sequence for 1A-3905  
SEQ ID NO: 73 is the determined cDNA sequence for V1-3686  
SEQ ID NO: 74 is the determined cDNA sequence for R1-2330  
SEQ ID NO: 75 is the determined cDNA sequence for 1B-3976  
SEQ ID NO: 76 is the determined cDNA sequence for V1-3679

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SEQ ID NO: 82 is the determined cDNA sequence for 1H-4774  
SEQ ID NO: 83 is the determined cDNA sequence for 1H-4781  
SEQ ID NO: 84 is the determined cDNA sequence for 1H-4785  
SEQ ID NO: 85 is the determined cDNA sequence for 1H-4787  
SEQ ID NO: 86 is the determined cDNA sequence for 1H-4796  
SEQ ID NO: 87 is the determined cDNA sequence for 1I-4807  
SEQ ID NO: 88 is the determined cDNA sequence for 1I-4810  
SEQ ID NO: 89 is the determined cDNA sequence for 1I-4811  
SEQ ID NO: 90 is the determined cDNA sequence for 1J-4876  
SEQ ID NO: 91 is the determined cDNA sequence for 1K-4884  
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SEQ ID NO: 107 is the determined full length cDNA sequence for F1-12 (also referred to as P504S)  
SEQ ID NO: 108 is the predicted amino acid sequence for F1-12  
SEQ ID NO: 109 is the determined full length cDNA sequence for J1-17  
SEQ ID NO: 110 is the determined full length cDNA sequence for L1-12  
SEQ ID NO: 111 is the determined full length cDNA sequence for N1-1862  
SEQ ID NO: 112 is the predicted amino acid sequence for J1-17



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SEQ ID NO: 202 is the determined extended cDNA sequence for 1D-4288  
SEQ ID NO: 203 is the determined extended cDNA sequence for 1D-4283  
SEQ ID NO: 204 is the determined extended cDNA sequence for 1D-4304  
SEQ ID NO: 205 is the determined extended cDNA sequence for 1D-4296  
SEQ ID NO: 206 is the determined extended cDNA sequence for 1D-4280  
SEQ ID NO: 207 is the determined cDNA sequence for 10-d8fwd  
SEQ ID NO: 208 is the determined cDNA sequence for 10-H10con  
SEQ ID NO: 209 is the determined cDNA sequence for 11-C8rev  
SEQ ID NO: 210 is the determined cDNA sequence for 7.g6fwd  
SEQ ID NO: 211 is the determined cDNA sequence for 7.g6rev  
SEQ ID NO: 212 is the determined cDNA sequence for 8-b5fwd  
SEQ ID NO: 213 is the determined cDNA sequence for 8-b5rev  
SEQ ID NO: 214 is the determined cDNA sequence for 8-b6fwd  
SEQ ID NO: 215 is the determined cDNA sequence for 8-b6 rev  
SEQ ID NO: 216 is the determined cDNA sequence for 8-d4fwd  
SEQ ID NO: 217 is the determined cDNA sequence for 8-d9rev  
SEQ ID NO: 218 is the determined cDNA sequence for 8-g3fwd  
SEQ ID NO: 219 is the determined cDNA sequence for 8-g3rev  
SEQ ID NO: 220 is the determined cDNA sequence for 8-h11rev  
SEQ ID NO: 221 is the determined cDNA sequence for g-f12fwd  
SEQ ID NO: 222 is the determined cDNA sequence for g-f3rev  
SEQ ID NO: 223 is the determined cDNA sequence for P509S

SEQ ID NO: 224 is the determined cDNA sequence for P510S  
SEQ ID NO: 225 is the determined cDNA sequence for P703DE5  
SEQ ID NO: 226 is the determined cDNA sequence for 9-A11  
SEQ ID NO: 227 is the determined cDNA sequence for 8-C6  
SEQ ID NO: 228 is the determined cDNA sequence for 8-H7  
SEQ ID NO: 229 is the determined cDNA sequence for JPTPN13  
SEQ ID NO: 230 is the determined cDNA sequence for JPTPN14  
SEQ ID NO: 231 is the determined cDNA sequence for JPTPN23  
SEQ ID NO: 232 is the determined cDNA sequence for JPTPN24  
SEQ ID NO: 233 is the determined cDNA sequence for JPTPN25  
SEQ ID NO: 234 is the determined cDNA sequence for JPTPN30  
SEQ ID NO: 235 is the determined cDNA sequence for JPTPN34  
SEQ ID NO: 236 is the determined cDNA sequence for PTPN35  
SEQ ID NO: 237 is the determined cDNA sequence for JPTPN36  
SEQ ID NO: 238 is the determined cDNA sequence for JPTPN38  
SEQ ID NO: 239 is the determined cDNA sequence for JPTPN39  
SEQ ID NO: 240 is the determined cDNA sequence for JPTPN40  
SEQ ID NO: 241 is the determined cDNA sequence for JPTPN41  
SEQ ID NO: 242 is the determined cDNA sequence for JPTPN42  
SEQ ID NO: 243 is the determined cDNA sequence for JPTPN45  
SEQ ID NO: 244 is the determined cDNA sequence for JPTPN46  
SEQ ID NO: 245 is the determined cDNA sequence for JPTPN51  
SEQ ID NO: 246 is the determined cDNA sequence for JPTPN56  
SEQ ID NO: 247 is the determined cDNA sequence for PTPN64  
SEQ ID NO: 248 is the determined cDNA sequence for JPTPN65  
SEQ ID NO: 249 is the determined cDNA sequence for JPTPN67  
SEQ ID NO: 250 is the determined cDNA sequence for JPTPN76  
SEQ ID NO: 251 is the determined cDNA sequence for JPTPN84  
SEQ ID NO: 252 is the determined cDNA sequence for JPTPN85  
SEQ ID NO: 253 is the determined cDNA sequence for JPTPN86  
SEQ ID NO: 254 is the determined cDNA sequence for JPTPN87  
SEQ ID NO: 255 is the determined cDNA sequence for JPTPN88  
SEQ ID NO: 256 is the determined cDNA sequence for JP1F1  
SEQ ID NO: 257 is the determined cDNA sequence for JP1F2  
SEQ ID NO: 258 is the determined cDNA sequence for JP1C2  
SEQ ID NO: 259 is the determined cDNA sequence for JP1B1  
SEQ ID NO: 260 is the determined cDNA sequence for JP1B2

SEQ ID NO: 261 is the determined cDNA sequence for JP1D3  
SEQ ID NO: 262 is the determined cDNA sequence for JP1A4  
SEQ ID NO: 263 is the determined cDNA sequence for JP1F5  
SEQ ID NO: 264 is the determined cDNA sequence for JP1E6  
SEQ ID NO: 265 is the determined cDNA sequence for JP1D6  
SEQ ID NO: 266 is the determined cDNA sequence for JP1B5  
SEQ ID NO: 267 is the determined cDNA sequence for JP1A6  
SEQ ID NO: 268 is the determined cDNA sequence for JP1E8  
SEQ ID NO: 269 is the determined cDNA sequence for JP1D7  
SEQ ID NO: 270 is the determined cDNA sequence for JP1D9  
SEQ ID NO: 271 is the determined cDNA sequence for JP1C10  
SEQ ID NO: 272 is the determined cDNA sequence for JP1A9  
SEQ ID NO: 273 is the determined cDNA sequence for JP1F12  
SEQ ID NO: 274 is the determined cDNA sequence for JP1E12  
SEQ ID NO: 275 is the determined cDNA sequence for JP1D11  
SEQ ID NO: 276 is the determined cDNA sequence for JP1C11  
SEQ ID NO: 277 is the determined cDNA sequence for JP1C12  
SEQ ID NO: 278 is the determined cDNA sequence for JP1B12  
SEQ ID NO: 279 is the determined cDNA sequence for JP1A12  
SEQ ID NO: 280 is the determined cDNA sequence for JP8G2  
SEQ ID NO: 281 is the determined cDNA sequence for JP8H1  
SEQ ID NO: 282 is the determined cDNA sequence for JP8H2  
SEQ ID NO: 283 is the determined cDNA sequence for JP8A3  
SEQ ID NO: 284 is the determined cDNA sequence for JP8A4  
SEQ ID NO: 285 is the determined cDNA sequence for JP8C3  
SEQ ID NO: 286 is the determined cDNA sequence for JP8G4  
SEQ ID NO: 287 is the determined cDNA sequence for JP8B6  
SEQ ID NO: 288 is the determined cDNA sequence for JP8D6  
SEQ ID NO: 289 is the determined cDNA sequence for JP8F5  
SEQ ID NO: 290 is the determined cDNA sequence for JP8A8  
SEQ ID NO: 291 is the determined cDNA sequence for JP8C7  
SEQ ID NO: 292 is the determined cDNA sequence for JP8D7  
SEQ ID NO: 293 is the determined cDNA sequence for P8D8  
SEQ ID NO: 294 is the determined cDNA sequence for JP8E7  
SEQ ID NO: 295 is the determined cDNA sequence for JP8F8  
SEQ ID NO: 296 is the determined cDNA sequence for JP8G8  
SEQ ID NO: 297 is the determined cDNA sequence for JP8B10

SEQ ID NO: 298 is the determined cDNA sequence for JP8C10  
SEQ ID NO: 299 is the determined cDNA sequence for JP8E9  
SEQ ID NO: 300 is the determined cDNA sequence for JP8E10  
SEQ ID NO: 301 is the determined cDNA sequence for JP8F9  
SEQ ID NO: 302 is the determined cDNA sequence for JP8H9  
SEQ ID NO: 303 is the determined cDNA sequence for JP8C12  
SEQ ID NO: 304 is the determined cDNA sequence for JP8E11  
SEQ ID NO: 305 is the determined cDNA sequence for JP8E12  
SEQ ID NO: 306 is the amino acid sequence for the peptide PS2#12  
SEQ ID NO: 307 is the determined cDNA sequence for P711P  
SEQ ID NO: 308 is the determined cDNA sequence for P712P  
SEQ ID NO: 309 is the determined cDNA sequence for CLONE23  
SEQ ID NO: 310 is the determined cDNA sequence for P774P  
SEQ ID NO: 311 is the determined cDNA sequence for P775P  
SEQ ID NO: 312 is the determined cDNA sequence for P715P  
SEQ ID NO: 313 is the determined cDNA sequence for P710P  
SEQ ID NO: 314 is the determined cDNA sequence for P767P  
SEQ ID NO: 315 is the determined cDNA sequence for P768P  
SEQ ID NO: 316-325 are the determined cDNA sequences of previously isolated genes  
SEQ ID NO: 326 is the determined cDNA sequence for P703PDE5  
SEQ ID NO: 327 is the predicted amino acid sequence for P703PDE5  
SEQ ID NO: 328 is the determined cDNA sequence for P703P6.26  
SEQ ID NO: 329 is the predicted amino acid sequence for P703P6.26  
SEQ ID NO: 330 is the determined cDNA sequence for P703PX-23  
SEQ ID NO: 331 is the predicted amino acid sequence for P703PX-23  
SEQ ID NO: 332 is the determined full length cDNA sequence for P509S  
SEQ ID NO: 333 is the determined extended cDNA sequence for P707P (also referred to as 11-C9)  
SEQ ID NO: 334 is the determined cDNA sequence for P714P  
SEQ ID NO: 335 is the determined cDNA sequence for P705P (also referred to as 9-F3)  
SEQ ID NO: 336 is the predicted amino acid sequence for P705P  
SEQ ID NO: 337 is the amino acid sequence of the peptide P1S#10  
SEQ ID NO: 338 is the amino acid sequence of the peptide p5  
SEQ ID NO: 339 is the predicted amino acid sequence of P509S  
SEQ ID NO: 340 is the determined cDNA sequence for P778P  
SEQ ID NO: 341 is the determined cDNA sequence for P786P  
SEQ ID NO: 342 is the determined cDNA sequence for P789P

SEQ ID NO: 343 is the determined cDNA sequence for a clone showing homology to Homo sapiens MM46 mRNA

SEQ ID NO: 344 is the determined cDNA sequence for a clone showing homology to Homo sapiens TNF-alpha stimulated ABC protein (ABC50) mRNA

SEQ ID NO: 345 is the determined cDNA sequence for a clone showing homology to Homo sapiens mRNA for E-cadherin

SEQ ID NO: 346 is the determined cDNA sequence for a clone showing homology to Human nuclear-encoded mitochondrial serine hydroxymethyltransferase (SHMT)

SEQ ID NO: 347 is the determined cDNA sequence for a clone showing homology to Homo sapiens natural resistance-associated macrophage protein2 (NRAMP2)

SEQ ID NO: 348 is the determined cDNA sequence for a clone showing homology to Homo sapiens phosphoglucomutase-related protein (PGMRP)

SEQ ID NO: 349 is the determined cDNA sequence for a clone showing homology to Human mRNA for proteosome subunit p40

SEQ ID NO: 350 is the determined cDNA sequence for P777P

SEQ ID NO: 351 is the determined cDNA sequence for P779P

SEQ ID NO: 352 is the determined cDNA sequence for P790P

SEQ ID NO: 353 is the determined cDNA sequence for P784P

SEQ ID NO: 354 is the determined cDNA sequence for P776P

SEQ ID NO: 355 is the determined cDNA sequence for P780P

SEQ ID NO: 356 is the determined cDNA sequence for P544S

SEQ ID NO: 357 is the determined cDNA sequence for P745S

SEQ ID NO: 358 is the determined cDNA sequence for P782P

SEQ ID NO: 359 is the determined cDNA sequence for P783P

SEQ ID NO: 360 is the determined cDNA sequence for unknown 17984

SEQ ID NO: 361 is the determined cDNA sequence for P787P

SEQ ID NO: 362 is the determined cDNA sequence for P788P

SEQ ID NO: 363 is the determined cDNA sequence for unknown 17994

SEQ ID NO: 364 is the determined cDNA sequence for P781P

SEQ ID NO: 365 is the determined cDNA sequence for P785P

SEQ ID NO: 366-375 are the determined cDNA sequences for splice variants of B305D.

SEQ ID NO: 376 is the predicted amino acid sequence encoded by the sequence of SEQ ID NO: 366.

SEQ ID NO: 377 is the predicted amino acid sequence encoded by the sequence of SEQ ID NO: 372.

SEQ ID NO: 378 is the predicted amino acid sequence encoded by the sequence of SEQ ID NO: 373.

SEQ ID NO: 379 is the predicted amino acid sequence encoded by the sequence of SEQ ID NO: 374.

SEQ ID NO: 380 is the predicted amino acid sequence encoded by the sequence of SEQ ID NO: 375.

SEQ ID NO: 381 is the determined cDNA sequence for B716P.

SEQ ID NO: 382 is the determined full-length cDNA sequence for P711P.

SEQ ID NO: 383 is the predicted amino acid sequence for P711P.

SEQ ID NO: 384 is the cDNA sequence for P1000C.

SEQ ID NO: 385 is the cDNA sequence for CGI-82.

SEQ ID NO: 386 is the cDNA sequence for 23320.

SEQ ID NO: 387 is the cDNA sequence for CGI-69.

SEQ ID NO: 388 is the cDNA sequence for L-iditol-2-dehydrogenase.

SEQ ID NO: 389 is the cDNA sequence for 23379.

SEQ ID NO: 390 is the cDNA sequence for 23381.

SEQ ID NO: 391 is the cDNA sequence for KIAA0122.

SEQ ID NO: 392 is the cDNA sequence for 23399.

SEQ ID NO: 393 is the cDNA sequence for a previously identified gene.

SEQ ID NO: 394 is the cDNA sequence for HCLBP.

SEQ ID NO: 395 is the cDNA sequence for transglutaminase.

SEQ ID NO: 396 is the cDNA sequence for a previously identified gene.

SEQ ID NO: 397 is the cDNA sequence for PAP.

SEQ ID NO: 398 is the cDNA sequence for Ets transcription factor PDEF.

SEQ ID NO: 399 is the cDNA sequence for hTGR.

SEQ ID NO: 400 is the cDNA sequence for KIAA0295.

SEQ ID NO: 401 is the cDNA sequence for 22545.

SEQ ID NO: 402 is the cDNA sequence for 22547.

SEQ ID NO: 403 is the cDNA sequence for 22548.

SEQ ID NO: 404 is the cDNA sequence for 22550.

SEQ ID NO: 405 is the cDNA sequence for 22551.

SEQ ID NO: 406 is the cDNA sequence for 22552.

SEQ ID NO: 407 is the cDNA sequence for 22553.

SEQ ID NO: 408 is the cDNA sequence for 22558.

SEQ ID NO: 409 is the cDNA sequence for 22562.

SEQ ID NO: 410 is the cDNA sequence for 22565.

SEQ ID NO: 411 is the cDNA sequence for 22567.

SEQ ID NO: 412 is the cDNA sequence for 22568.

SEQ ID NO: 413 is the cDNA sequence for 22570.



SEQ ID NO:414 is the cDNA sequence for 22571.  
SEQ ID NO:415 is the cDNA sequence for 22572.  
SEQ ID NO:416 is the cDNA sequence for 22573.  
SEQ ID NO:417 is the cDNA sequence for 22573.  
SEQ ID NO:418 is the cDNA sequence for 22575.  
SEQ ID NO:419 is the cDNA sequence for 22580.  
SEQ ID NO:420 is the cDNA sequence for 22581.  
SEQ ID NO:421 is the cDNA sequence for 22582.  
SEQ ID NO:422 is the cDNA sequence for 22583.  
SEQ ID NO:423 is the cDNA sequence for 22584.  
SEQ ID NO:424 is the cDNA sequence for 22585.  
SEQ ID NO:425 is the cDNA sequence for 22586.  
SEQ ID NO:426 is the cDNA sequence for 22587.  
SEQ ID NO:427 is the cDNA sequence for 22588.  
SEQ ID NO:428 is the cDNA sequence for 22589.  
SEQ ID NO:429 is the cDNA sequence for 22590.  
SEQ ID NO:430 is the cDNA sequence for 22591.  
SEQ ID NO:431 is the cDNA sequence for 22592.  
SEQ ID NO:432 is the cDNA sequence for 22593.  
SEQ ID NO:433 is the cDNA sequence for 22594.  
SEQ ID NO:434 is the cDNA sequence for 22595.  
SEQ ID NO:435 is the cDNA sequence for 22596.  
SEQ ID NO:436 is the cDNA sequence for 22847.  
SEQ ID NO:437 is the cDNA sequence for 22848.  
SEQ ID NO:438 is the cDNA sequence for 22849.  
SEQ ID NO:439 is the cDNA sequence for 22851.  
SEQ ID NO:440 is the cDNA sequence for 22852.  
SEQ ID NO:441 is the cDNA sequence for 22853.  
SEQ ID NO:442 is the cDNA sequence for 22854.  
SEQ ID NO:443 is the cDNA sequence for 22855.  
SEQ ID NO:444 is the cDNA sequence for 22856.  
SEQ ID NO:445 is the cDNA sequence for 22857.  
SEQ ID NO:446 is the cDNA sequence for 23601.  
SEQ ID NO:447 is the cDNA sequence for 23602.  
SEQ ID NO:448 is the cDNA sequence for 23605.  
SEQ ID NO:449 is the cDNA sequence for 23606.  
SEQ ID NO:450 is the cDNA sequence for 23612.

SEQ ID NO:451 is the cDNA sequence for 23614.

SEQ ID NO:452 is the cDNA sequence for 23618.

SEQ ID NO:453 is the cDNA sequence for 23622.

SEQ ID NO:454 is the cDNA sequence for folate hydrolase.

SEQ ID NO:455 is the cDNA sequence for LIM protein.

SEQ ID NO:456 is the cDNA sequence for a known gene.

SEQ ID NO:457 is the cDNA sequence for a known gene.

SEQ ID NO:458 is the cDNA sequence for a previously identified gene.

SEQ ID NO:459 is the cDNA sequence for 23045.

SEQ ID NO:460 is the cDNA sequence for 23032.

SEQ ID NO:461 is the cDNA sequence for 23054.

SEQ ID NOs:462-467 are cDNA sequences for known genes.

SEQ ID NOs:468-471 are cDNA sequences for P710P.

SEQ ID NO:472 is a cDNA sequence for P1001C.

#### DETAILED DESCRIPTION OF THE INVENTION

As noted above, the present invention is generally directed to compositions and methods for the therapy and diagnosis of cancer, such as prostate cancer. The compositions described herein may include prostate tumor polypeptides, polynucleotides encoding such polypeptides, binding agents such as antibodies, antigen presenting cells (APCs) and/or immune system cells (*e.g.*, T cells). Polypeptides of the present invention generally comprise at least a portion (such as an immunogenic portion) of a prostate tumor protein or a variant thereof. A "prostate tumor protein" is a protein that is expressed in prostate tumor cells at a level that is at least two fold, and preferably at least five fold, greater than the level of expression in a normal tissue, as determined using a representative assay provided herein. Certain prostate tumor proteins are tumor proteins that react detectably (within an immunoassay, such as an ELISA or Western blot) with antisera of a patient afflicted with prostate cancer. Polynucleotides of the subject invention generally comprise a DNA or RNA sequence that encodes all or a portion of such a polypeptide, or that is complementary to such a sequence. Antibodies are generally immune system proteins, or antigen-binding fragments thereof, that are capable of binding to a polypeptide as described above. Antigen presenting cells include dendritic cells, macrophages, monocytes, fibroblasts and B-cells that express a polypeptide as described above. T cells that may be employed within such compositions are generally T cells that are specific for a polypeptide as described above.

The present invention is based on the discovery of human prostate tumor proteins. Sequences of polynucleotides encoding certain tumor proteins, or portions thereof, are provided in SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472. Sequences of polypeptides comprising at least a portion of a tumor protein are provided in SEQ ID NOs:112-114, 172, 176, 178, 327, 329, 331, 336, 339, 376-380 and 383.

#### PROSTATE TUMOR PROTEIN POLYNUCLEOTIDES

Any polynucleotide that encodes a prostate tumor protein or a portion or other variant thereof as described herein is encompassed by the present invention. Preferred polynucleotides comprise at least 15 consecutive nucleotides, preferably at least 30 consecutive nucleotides and more preferably at least 45 consecutive nucleotides, that encode a portion of a prostate tumor protein. More preferably, a polynucleotide encodes an immunogenic portion of a prostate tumor protein. Polynucleotides complementary to any such sequences are also encompassed by the present invention. Polynucleotides may be single-stranded (coding or antisense) or double-stranded, and may be DNA (genomic, cDNA or synthetic) or RNA molecules. RNA molecules include HnRNA molecules, which contain introns and correspond to a DNA molecule in a one-to-one manner, and mRNA molecules, which do not contain introns. Additional coding or non-coding sequences may, but need not, be present within a polynucleotide of the present invention, and a polynucleotide may, but need not, be linked to other molecules and/or support materials.

Polynucleotides may comprise a native sequence (*i.e.*, an endogenous sequence that encodes a prostate tumor protein or a portion thereof) or may comprise a variant of such a sequence. Polynucleotide variants may contain one or more substitutions, additions, deletions and/or insertions such that the immunogenicity of the encoded polypeptide is not diminished, relative to a native tumor protein. The effect on the immunogenicity of the encoded polypeptide may generally be assessed as described herein. Variants preferably exhibit at least about 70% identity, more preferably at least about 80% identity and most preferably at least about 90% identity to a polynucleotide sequence that encodes a native prostate tumor protein or a portion thereof.

Two polynucleotide or polypeptide sequences are said to be "identical" if the sequence of nucleotides or amino acids in the two sequences is the same when aligned for maximum correspondence as described below. Comparisons between two sequences are typically performed by comparing the sequences over a comparison window to identify and compare local regions of sequence similarity. A "comparison window" as used herein, refers to a segment of at least about 20 contiguous positions, usually 30 to about 75, 40 to about 50,

in which a sequence may be compared to a reference sequence of the same number of contiguous positions after the two sequences are optimally aligned.

Optimal alignment of sequences for comparison may be conducted using the Megalign program in the Lasergene suite of bioinformatics software (DNASTAR, Inc., Madison, WI), using default parameters. This program embodies several alignment schemes described in the following references: Dayhoff, M.O. (1978) A model of evolutionary change in proteins – Matrices for detecting distant relationships. In Dayhoff, M.O. (ed.) *Atlas of Protein Sequence and Structure*, National Biomedical Research Foundation, Washington DC Vol. 5, Suppl. 3, pp. 345-358; Hein J. (1990) Unified Approach to Alignment and Phylogenesis pp. 626-645 *Methods in Enzymology* vol. 183, Academic Press, Inc., San Diego, CA; Higgins, D.G. and Sharp, P.M. (1989) *CABIOS* 5:151-153; Myers, E.W. and Muller W. (1988) *CABIOS* 4:11-17; Robinson, E.D. (1971) *Comb. Theor* 11:105; Santou, N. Nes, M. (1987) *Mol. Biol. Evol.* 4:406-425; Sneath, P.H.A. and Sokal, R.R. (1973) *Numerical Taxonomy – the Principles and Practice of Numerical Taxonomy*, Freeman Press, San Francisco, CA; Wilbur, W.J. and Lipman, D.J. (1983) *Proc. Natl. Acad., Sci. USA* 80:726-730.

Preferably, the “percentage of sequence identity” is determined by comparing two optimally aligned sequences over a window of comparison of at least 20 positions, wherein the portion of the polynucleotide or polypeptide sequence in the comparison window may comprise additions or deletions (*i.e.*, gaps) of 20 percent or less, usually 5 to 15 percent, or 10 to 12 percent, as compared to the reference sequences (which does not comprise additions or deletions) for optimal alignment of the two sequences. The percentage is calculated by determining the number of positions at which the identical nucleic acid bases or amino acid residue occurs in both sequences to yield the number of matched positions, dividing the number of matched positions by the total number of positions in the reference sequence (*i.e.*, the window size) and multiplying the results by 100 to yield the percentage of sequence identity.

Variants may also, or alternatively, be substantially homologous to a native gene, or a portion or complement thereof. Such polynucleotide variants are capable of hybridizing under moderately stringent conditions to a naturally occurring DNA sequence encoding a native prostate tumor protein (or a complementary sequence). Suitable moderately stringent conditions include prewashing in a solution of 5 X SSC, 0.5% SDS, 1.0 mM EDTA (pH 8.0); hybridizing at 50°C-65°C, 5 X SSC; overnight; followed by washing twice at 65°C for 20 minutes with each of 2X, 0.5X and 0.2X SSC containing 0.1% SDS.

It will be appreciated by those of ordinary skill in the art that, as a result of the degeneracy of the genetic code, there are many nucleotide sequences that encode a polypeptide as described herein. Some of these polynucleotides bear minimal homology to

the nucleotide sequence of any native gene. Nonetheless, polynucleotides that vary due to differences in codon usage are specifically contemplated by the present invention. Further, alleles of the genes comprising the polynucleotide sequences provided herein are within the scope of the present invention. Alleles are endogenous genes that are altered as a result of one or more mutations, such as deletions, additions and/or substitutions of nucleotides. The resulting mRNA and protein may, but need not, have an altered structure or function. Alleles may be identified using standard techniques (such as hybridization, amplification and/or database sequence comparison).

Polynucleotides may be prepared using any of a variety of techniques. For example, a polynucleotide may be identified, as described in more detail below, by screening a microarray of cDNAs for tumor-associated expression (*i.e.*, expression that is at least five fold greater in a prostate tumor than in normal tissue, as determined using a representative assay provided herein). Such screens may be performed using a Synteni microarray (Palo Alto, CA) according to the manufacturer's instructions (and essentially as described by Schena et al., *Proc. Natl. Acad. Sci. USA* 93:10614-10619, 1996 and Heller et al., *Proc. Natl. Acad. Sci. USA* 94:2150-2155, 1997). Alternatively, polypeptides may be amplified from cDNA prepared from cells expressing the proteins described herein, such as prostate tumor cells. Such polynucleotides may be amplified via polymerase chain reaction (PCR). For this approach, sequence-specific primers may be designed based on the sequences provided herein, and may be purchased or synthesized.

An amplified portion may be used to isolate a full length gene from a suitable library (*e.g.*, a prostate tumor cDNA library) using well known techniques. Within such techniques, a library (cDNA or genomic) is screened using one or more polynucleotide probes or primers suitable for amplification. Preferably, a library is size-selected to include larger molecules. Random primed libraries may also be preferred for identifying 5' and upstream regions of genes. Genomic libraries are preferred for obtaining introns and extending 5' sequences.

For hybridization techniques, a partial sequence may be labeled (*e.g.*, by nick-translation or end-labeling with  $^{32}\text{P}$ ) using well known techniques. A bacterial or bacteriophage library is then screened by hybridizing filters containing denatured bacterial colonies (or lawns containing phage plaques) with the labeled probe (*see* Sambrook et al., *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY, 1989). Hybridizing colonies or plaques are selected and expanded, and the DNA is isolated for further analysis. cDNA clones may be analyzed to determine the amount of additional sequence by, for example, PCR using a primer from the partial sequence and a primer from the vector. Restriction maps and partial sequences may be generated to identify one or more overlapping clones. The complete sequence may then be determined using

standard techniques, which may involve generating a series of deletion clones. The resulting overlapping sequences are then assembled into a single contiguous sequence. A full length cDNA molecule can be generated by ligating suitable fragments, using well known techniques.

Alternatively, there are numerous amplification techniques for obtaining a full length coding sequence from a partial cDNA sequence. Within such techniques, amplification is generally performed via PCR. Any of a variety of commercially available kits may be used to perform the amplification step. Primers may be designed using, for example, software well known in the art. Primers are preferably 22-30 nucleotides in length, have a GC content of at least 50% and anneal to the target sequence at temperatures of about 68°C to 72°C. The amplified region may be sequenced as described above, and overlapping sequences assembled into a contiguous sequence.

One such amplification technique is inverse PCR (*see* Triglia et al., *Nucl. Acids Res.* 16:8186, 1988), which uses restriction enzymes to generate a fragment in the known region of the gene. The fragment is then circularized by intramolecular ligation and used as a template for PCR with divergent primers derived from the known region. Within an alternative approach, sequences adjacent to a partial sequence may be retrieved by amplification with a primer to a linker sequence and a primer specific to a known region. The amplified sequences are typically subjected to a second round of amplification with the same linker primer and a second primer specific to the known region. A variation on this procedure, which employs two primers that initiate extension in opposite directions from the known sequence, is described in WO 96/38591. Another such technique is known as "rapid amplification of cDNA ends" or RACE. This technique involves the use of an internal primer and an external primer, which hybridizes to a polyA region or vector sequence, to identify sequences that are 5' and 3' of a known sequence. Additional techniques include capture PCR (Lagerstrom et al., *PCR Methods Applic.* 1:111-19, 1991) and walking PCR (Parker et al., *Nucl. Acids Res.* 19:3055-60, 1991). Other methods employing amplification may also be employed to obtain a full length cDNA sequence.

In certain instances, it is possible to obtain a full length cDNA sequence by analysis of sequences provided in an expressed sequence tag (EST) database, such as that available from GenBank. Searches for overlapping ESTs may generally be performed using well known programs (*e.g.*, NCBI BLAST searches), and such ESTs may be used to generate a contiguous full length sequence.

Certain nucleic acid sequences of cDNA molecules encoding at least a portion of a prostate tumor protein are provided in SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472. Isolation of these

polynucleotides is described below. Each of these prostate tumor proteins was overexpressed in prostate tumor tissue.

Polynucleotide variants may generally be prepared by any method known in the art, including chemical synthesis by, for example, solid phase phosphoramidite chemical synthesis. Modifications in a polynucleotide sequence may also be introduced using standard mutagenesis techniques, such as oligonucleotide-directed site-specific mutagenesis (*see* Adelman et al., *DNA* 2:183, 1983). Alternatively, RNA molecules may be generated by *in vitro* or *in vivo* transcription of DNA sequences encoding a prostate tumor protein, or portion thereof, provided that the DNA is incorporated into a vector with a suitable RNA polymerase promoter (such as T7 or SP6). Certain portions may be used to prepare an encoded polypeptide, as described herein. In addition, or alternatively, a portion may be administered to a patient such that the encoded polypeptide is generated *in vivo* (e.g., by transfecting antigen-presenting cells, such as dendritic cells, with a cDNA construct encoding a prostate tumor polypeptide, and administering the transfected cells to the patient).

A portion of a sequence complementary to a coding sequence (*i.e.*, an antisense polynucleotide) may also be used as a probe or to modulate gene expression. cDNA constructs that can be transcribed into antisense RNA may also be introduced into cells of tissues to facilitate the production of antisense RNA. An antisense polynucleotide may be used, as described herein, to inhibit expression of a tumor protein. Antisense technology can be used to control gene expression through triple-helix formation, which compromises the ability of the double helix to open sufficiently for the binding of polymerases, transcription factors or regulatory molecules (*see* Gee et al., *In Huber and Carr, Molecular and Immunologic Approaches*, Futura Publishing Co. (Mt. Kisco, NY; 1994)). Alternatively, an antisense molecule may be designed to hybridize with a control region of a gene (e.g., promoter, enhancer or transcription initiation site), and block transcription of the gene; or to block translation by inhibiting binding of a transcript to ribosomes.

A portion of a coding sequence, or of a complementary sequence, may also be designed as a probe or primer to detect gene expression. Probes may be labeled with a variety of reporter groups, such as radionuclides and enzymes, and are preferably at least 10 nucleotides in length, more preferably at least 20 nucleotides in length and still more preferably at least 30 nucleotides in length. Primers, as noted above, are preferably 22-30 nucleotides in length.

Any polynucleotide may be further modified to increase stability *in vivo*. Possible modifications include, but are not limited to, the addition of flanking sequences at the 5' and/or 3' ends; the use of phosphorothioate or 2' O-methyl rather than phosphodiesterase linkages in the backbone; and/or the inclusion of nontraditional bases such

as inosine, queosine and wybutosine, as well as acetyl-, methyl-, thio- and other modified forms of adenine, cytidine, guanine, thymine and uridine.

Nucleotide sequences as described herein may be joined to a variety of other nucleotide sequences using established recombinant DNA techniques. For example, a polynucleotide may be cloned into any of a variety of cloning vectors, including plasmids, phagemids, lambda phage derivatives and cosmids. Vectors of particular interest include expression vectors, replication vectors, probe generation vectors and sequencing vectors. In general, a vector will contain an origin of replication functional in at least one organism, convenient restriction endonuclease sites and one or more selectable markers. Other elements will depend upon the desired use, and will be apparent to those of ordinary skill in the art.

Within certain embodiments, polynucleotides may be formulated so as to permit entry into a cell of a mammal, and expression therein. Such formulations are particularly useful for therapeutic purposes, as described below. Those of ordinary skill in the art will appreciate that there are many ways to achieve expression of a polynucleotide in a target cell, and any suitable method may be employed. For example, a polynucleotide may be incorporated into a viral vector such as, but not limited to, adenovirus, adeno-associated virus, retrovirus, or vaccinia or other pox virus (*e.g.*, avian pox virus). Techniques for incorporating DNA into such vectors are well known to those of ordinary skill in the art. A retroviral vector may additionally transfer or incorporate a gene for a selectable marker (to aid in the identification or selection of transduced cells) and/or a targeting moiety, such as a gene that encodes a ligand for a receptor on a specific target cell, to render the vector target specific. Targeting may also be accomplished using an antibody, by methods known to those of ordinary skill in the art.

Other formulations for therapeutic purposes include colloidal dispersion systems, such as macromolecule complexes, nanocapsules, microspheres, beads, and lipid-based systems including oil-in-water emulsions, micelles, mixed micelles, and liposomes. A preferred colloidal system for use as a delivery vehicle *in vitro* and *in vivo* is a liposome (*i.e.*, an artificial membrane vesicle). The preparation and use of such systems is well known in the art.

#### PROSTATE TUMOR POLYPEPTIDES

Within the context of the present invention, polypeptides may comprise at least an immunogenic portion of a prostate tumor protein or a variant thereof, as described herein. As noted above, a "prostate tumor protein" is a protein that is expressed by prostate tumor cells. Proteins that are prostate tumor proteins also react detectably within an immunoassay (such as an ELISA) with antisera from a patient with prostate cancer. Polypeptides as described herein may be of any length. Additional sequences derived from



the native protein and/or heterologous sequences may be present, and such sequences may (but need not) possess further immunogenic or antigenic properties.

An "immunogenic portion," as used herein is a portion of a protein that is recognized (*i.e.*, specifically bound) by a B-cell and/or T-cell surface antigen receptor. Such immunogenic portions generally comprise at least 5 amino acid residues, more preferably at least 10, and still more preferably at least 20 amino acid residues of a prostate tumor protein or a variant thereof. Certain preferred immunogenic portions include peptides in which an N-terminal leader sequence and/or transmembrane domain have been deleted. Other preferred immunogenic portions may contain a small N- and/or C-terminal deletion (*e.g.*, 1-30 amino acids, preferably 5-15 amino acids), relative to the mature protein.

Immunogenic portions may generally be identified using well known techniques, such as those summarized in Paul, *Fundamental Immunology*, 3rd ed., 243-247 (Raven Press, 1993) and references cited therein. Such techniques include screening polypeptides for the ability to react with antigen-specific antibodies, antisera and/or T-cell lines or clones. As used herein, antisera and antibodies are "antigen-specific" if they specifically bind to an antigen (*i.e.*, they react with the protein in an ELISA or other immunoassay, and do not react detectably with unrelated proteins). Such antisera and antibodies may be prepared as described herein, and using well known techniques. An immunogenic portion of a native prostate tumor protein is a portion that reacts with such antisera and/or T-cells at a level that is not substantially less than the reactivity of the full length polypeptide (*e.g.*, in an ELISA and/or T-cell reactivity assay). Such immunogenic portions may react within such assays at a level that is similar to or greater than the reactivity of the full length polypeptide. Such screens may generally be performed using methods well known to those of ordinary skill in the art, such as those described in Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. For example, a polypeptide may be immobilized on a solid support and contacted with patient sera to allow binding of antibodies within the sera to the immobilized polypeptide. Unbound sera may then be removed and bound antibodies detected using, for example, <sup>125</sup>I-labeled Protein A.

As noted above, a composition may comprise a variant of a native prostate tumor protein. A polypeptide "variant," as used herein, is a polypeptide that differs from a native prostate tumor protein in one or more substitutions, deletions, additions and/or insertions, such that the immunogenicity of the polypeptide is not substantially diminished. In other words, the ability of a variant to react with antigen-specific antisera may be enhanced or unchanged, relative to the native protein, or may be diminished by less than 50%, and preferably less than 20%, relative to the native protein. Such variants may generally be identified by modifying one of the above polypeptide sequences and evaluating the reactivity of the modified polypeptide with antigen-specific antibodies or antisera as described herein.

Preferred variants include those in which one or more portions, such as an N-terminal leader sequence or transmembrane domain, have been removed. Other preferred variants include variants in which a small portion (*e.g.*, 1-30 amino acids, preferably 5-15 amino acids) has been removed from the N- and/or C-terminal of the mature protein. Polypeptide variants preferably exhibit at least about 70%, more preferably at least about 90% and most preferably at least about 95% identity (determined as described above) to the identified polypeptides.

Preferably, a variant contains conservative substitutions. A "conservative substitution" is one in which an amino acid is substituted for another amino acid that has similar properties, such that one skilled in the art of peptide chemistry would expect the secondary structure and hydrophobic nature of the polypeptide to be substantially unchanged. Amino acid substitutions may generally be made on the basis of similarity in polarity, charge, solubility, hydrophobicity, hydrophilicity and/or the amphipathic nature of the residues. For example, negatively charged amino acids include aspartic acid and glutamic acid; positively charged amino acids include lysine and arginine; and amino acids with uncharged polar head groups having similar hydrophilicity values include leucine, isoleucine and valine; glycine and alanine; asparagine and glutamine; and serine, threonine, phenylalanine and tyrosine. Other groups of amino acids that may represent conservative changes include: (1) ala, pro, gly, glu, asp, gln, asn, ser, thr; (2) cys, ser, tyr, thr; (3) val, ile, leu, met, ala, phe; (4) lys, arg, his; and (5) phe, tyr, trp, his. A variant may also, or alternatively, contain nonconservative changes. In a preferred embodiment, variant polypeptides differ from a native sequence by substitution, deletion or addition of five amino acids or fewer. Variants may also (or alternatively) be modified by, for example, the deletion or addition of amino acids that have minimal influence on the immunogenicity, secondary structure and hydrophobic nature of the polypeptide.

As noted above, polypeptides may comprise a signal (or leader) sequence at the N-terminal end of the protein which co-translationally or post-translationally directs transfer of the protein. The polypeptide may also be conjugated to a linker or other sequence for ease of synthesis, purification or identification of the polypeptide (*e.g.*, poly-His), or to enhance binding of the polypeptide to a solid support. For example, a polypeptide may be conjugated to an immunoglobulin Fc region.

Polypeptides may be prepared using any of a variety of well known techniques. Recombinant polypeptides encoded by DNA sequences as described above may be readily prepared from the DNA sequences using any of a variety of expression vectors known to those of ordinary skill in the art. Expression may be achieved in any appropriate host cell that has been transformed or transfected with an expression vector containing a DNA molecule that encodes a recombinant polypeptide. Suitable host cells include prokaryotes, yeast and higher eukaryotic cells. Preferably, the host cells employed are

*E. coli*, yeast or a mammalian cell line such as COS or CHO. Supernatants from suitable host/vector systems which secrete recombinant protein or polypeptide into culture media may be first concentrated using a commercially available filter. Following concentration, the concentrate may be applied to a suitable purification matrix such as an affinity matrix or an ion exchange resin. Finally, one or more reverse phase HPLC steps can be employed to further purify a recombinant polypeptide.

Portions and other variants having fewer than about 100 amino acids, and generally fewer than about 50 amino acids, may also be generated by synthetic means, using techniques well known to those of ordinary skill in the art. For example, such polypeptides may be synthesized using any of the commercially available solid-phase techniques, such as the Merrifield solid-phase synthesis method, where amino acids are sequentially added to a growing amino acid chain. See Merrifield, *J. Am. Chem. Soc.* 85:2149-2146, 1963. Equipment for automated synthesis of polypeptides is commercially available from suppliers such as Perkin Elmer/Applied BioSystems Division (Foster City, CA), and may be operated according to the manufacturer's instructions.

Within certain specific embodiments, a polypeptide may be a fusion protein that comprises multiple polypeptides as described herein, or that comprises at least one polypeptide as described herein and an unrelated sequence, such as a known tumor protein. A fusion partner may, for example, assist in providing T helper epitopes (an immunological fusion partner), preferably T helper epitopes recognized by humans, or may assist in expressing the protein (an expression enhancer) at higher yields than the native recombinant protein. Certain preferred fusion partners are both immunological and expression enhancing fusion partners. Other fusion partners may be selected so as to increase the solubility of the protein or to enable the protein to be targeted to desired intracellular compartments. Still further fusion partners include affinity tags, which facilitate purification of the protein.

Fusion proteins may generally be prepared using standard techniques, including chemical conjugation. Preferably, a fusion protein is expressed as a recombinant protein, allowing the production of increased levels, relative to a non-fused protein, in an expression system. Briefly, DNA sequences encoding the polypeptide components may be assembled separately, and ligated into an appropriate expression vector. The 3' end of the DNA sequence encoding one polypeptide component is ligated, with or without a peptide linker, to the 5' end of a DNA sequence encoding the second polypeptide component so that the reading frames of the sequences are in phase. This permits translation into a single fusion protein that retains the biological activity of both component polypeptides.

A peptide linker sequence may be employed to separate the first and the second polypeptide components by a distance sufficient to ensure that each polypeptide folds into its secondary and tertiary structures. Such a peptide linker sequence is incorporated into

the fusion protein using standard techniques well known in the art. Suitable peptide linker sequences may be chosen based on the following factors: (1) their ability to adopt a flexible extended conformation; (2) their inability to adopt a secondary structure that could interact with functional epitopes on the first and second polypeptides; and (3) the lack of hydrophobic or charged residues that might react with the polypeptide functional epitopes. Preferred peptide linker sequences contain Gly, Asn and Ser residues. Other near neutral amino acids, such as Thr and Ala may also be used in the linker sequence. Amino acid sequences which may be usefully employed as linkers include those disclosed in Maratea et al., *Gene* 40:39-46, 1985; Murphy et al., *Proc. Natl. Acad. Sci. USA* 83:8258-8262, 1986; U.S. Patent No. 4,935,233 and U.S. Patent No. 4,751,180. The linker sequence may generally be from 1 to about 50 amino acids in length. Linker sequences are not required when the first and second polypeptides have non-essential N-terminal amino acid regions that can be used to separate the functional domains and prevent steric interference.

The ligated DNA sequences are operably linked to suitable transcriptional or translational regulatory elements. The regulatory elements responsible for expression of DNA are located only 5' to the DNA sequence encoding the first polypeptides. Similarly, stop codons required to end translation and transcription termination signals are only present 3' to the DNA sequence encoding the second polypeptide.

Fusion proteins are also provided that comprise a polypeptide of the present invention together with an unrelated immunogenic protein. Preferably the immunogenic protein is capable of eliciting a recall response. Examples of such proteins include tetanus, tuberculosis and hepatitis proteins (*see, for example, Stoute et al. New Engl. J. Med.*, 336:86-91, 1997).

Within preferred embodiments, an immunological fusion partner is derived from protein D, a surface protein of the gram-negative bacterium *Haemophilus influenza B* (WO 91/18926). Preferably, a protein D derivative comprises approximately the first third of the protein (*e.g.*, the first N-terminal 100-110 amino acids), and a protein D derivative may be lipidated. Within certain preferred embodiments, the first 109 residues of a Lipoprotein D fusion partner is included on the N-terminus to provide the polypeptide with additional exogenous T-cell epitopes and to increase the expression level in *E. coli* (thus functioning as an expression enhancer). The lipid tail ensures optimal presentation of the antigen to antigen presenting cells. Other fusion partners include the non-structural protein from influenzae virus, NS1 (hemagglutinin). Typically, the N-terminal 81 amino acids are used, although different fragments that include T-helper epitopes may be used.

In another embodiment, the immunological fusion partner is the protein known as LYTA, or a portion thereof (preferably a C-terminal portion). LYTA is derived from *Streptococcus pneumoniae*, which synthesizes an N-acetyl-L-alanine amidase known as

amidase LYTA (encoded by the *LytA* gene; *Gene* 43:265-292, 1986). LYTA is an autolysin that specifically degrades certain bonds in the peptidoglycan backbone. The C-terminal domain of the LYTA protein is responsible for the affinity to the choline or to some choline analogues such as DEAE. This property has been exploited for the development of *E. coli* C-LYTA expressing plasmids useful for expression of fusion proteins. Purification of hybrid proteins containing the C-LYTA fragment at the amino terminus has been described (*see Biotechnology* 10:795-798, 1992). Within a preferred embodiment, a repeat portion of LYTA may be incorporated into a fusion protein. A repeat portion is found in the C-terminal region starting at residue 178. A particularly preferred repeat portion incorporates residues 188-305.

In general, polypeptides (including fusion proteins) and polynucleotides as described herein are isolated. An "isolated" polypeptide or polynucleotide is one that is removed from its original environment. For example, a naturally-occurring protein is isolated if it is separated from some or all of the coexisting materials in the natural system. Preferably, such polypeptides are at least about 90% pure, more preferably at least about 95% pure and most preferably at least about 99% pure. A polynucleotide is considered to be isolated if, for example, it is cloned into a vector that is not a part of the natural environment.

#### BINDING AGENTS

The present invention further provides agents, such as antibodies and antigen-binding fragments thereof, that specifically bind to a prostate tumor protein. As used herein, an antibody, or antigen-binding fragment thereof, is said to "specifically bind" to a prostate tumor protein if it reacts at a detectable level (within, for example, an ELISA) with a prostate tumor protein, and does not react detectably with unrelated proteins under similar conditions. As used herein, "binding" refers to a noncovalent association between two separate molecules such that a complex is formed. The ability to bind may be evaluated by, for example, determining a binding constant for the formation of the complex. The binding constant is the value obtained when the concentration of the complex is divided by the product of the component concentrations. In general, two compounds are said to "bind," in the context of the present invention, when the binding constant for complex formation exceeds about  $10^3$  L/mol. The binding constant may be determined using methods well known in the art.

Binding agents may be further capable of differentiating between patients with and without a cancer, such as prostate cancer, using the representative assays provided herein. In other words, antibodies or other binding agents that bind to a prostate tumor protein will generate a signal indicating the presence of a cancer in at least about 20% of patients with the disease, and will generate a negative signal indicating the absence of the disease in at least about 90% of individuals without the cancer. To determine whether a binding agent satisfies this requirement, biological samples (*e.g.*, blood, sera, urine and/or tumor biopsies) from

patients with and without a cancer (as determined using standard clinical tests) may be assayed as described herein for the presence of polypeptides that bind to the binding agent. It will be apparent that a statistically significant number of samples with and without the disease should be assayed. Each binding agent should satisfy the above criteria; however, those of ordinary skill in the art will recognize that binding agents may be used in combination to improve sensitivity.

Any agent that satisfies the above requirements may be a binding agent. For example, a binding agent may be a ribosome, with or without a peptide component, an RNA molecule or a polypeptide. In a preferred embodiment, a binding agent is an antibody or an antigen-binding fragment thereof. Antibodies may be prepared by any of a variety of techniques known to those of ordinary skill in the art. *See, e.g., Harlow and Lane, Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. In general, antibodies can be produced by cell culture techniques, including the generation of monoclonal antibodies as described herein, or via transfection of antibody genes into suitable bacterial or mammalian cell hosts, in order to allow for the production of recombinant antibodies. In one technique, an immunogen comprising the polypeptide is initially injected into any of a wide variety of mammals (*e.g., mice, rats, rabbits, sheep or goats*). In this step, the polypeptides of this invention may serve as the immunogen without modification. Alternatively, particularly for relatively short polypeptides, a superior immune response may be elicited if the polypeptide is joined to a carrier protein, such as bovine serum albumin or keyhole limpet hemocyanin. The immunogen is injected into the animal host, preferably according to a predetermined schedule incorporating one or more booster immunizations, and the animals are bled periodically. Polyclonal antibodies specific for the polypeptide may then be purified from such antisera by, for example, affinity chromatography using the polypeptide coupled to a suitable solid support.

Monoclonal antibodies specific for an antigenic polypeptide of interest may be prepared, for example, using the technique of Kohler and Milstein, *Eur. J. Immunol.* 6:511-519, 1976, and improvements thereto. Briefly, these methods involve the preparation of immortal cell lines capable of producing antibodies having the desired specificity (*i.e., reactivity with the polypeptide of interest*). Such cell lines may be produced, for example, from spleen cells obtained from an animal immunized as described above. The spleen cells are then immortalized by, for example, fusion with a myeloma cell fusion partner, preferably one that is syngeneic with the immunized animal. A variety of fusion techniques may be employed. For example, the spleen cells and myeloma cells may be combined with a nonionic detergent for a few minutes and then plated at low density on a selective medium that supports the growth of hybrid cells, but not myeloma cells. A preferred selection technique uses HAT (hypoxanthine, aminopterin, thymidine) selection. After a sufficient

time, usually about 1 to 2 weeks, colonies of hybrids are observed. Single colonies are selected and their culture supernatants tested for binding activity against the polypeptide. Hybridomas having high reactivity and specificity are preferred.

Monoclonal antibodies may be isolated from the supernatants of growing hybridoma colonies. In addition, various techniques may be employed to enhance the yield, such as injection of the hybridoma cell line into the peritoneal cavity of a suitable vertebrate host, such as a mouse. Monoclonal antibodies may then be harvested from the ascites fluid or the blood. Contaminants may be removed from the antibodies by conventional techniques, such as chromatography, gel filtration, precipitation, and extraction. The polypeptides of this invention may be used in the purification process in, for example, an affinity chromatography step.

Within certain embodiments, the use of antigen-binding fragments of antibodies may be preferred. Such fragments include Fab fragments, which may be prepared using standard techniques. Briefly, immunoglobulins may be purified from rabbit serum by affinity chromatography on Protein A bead columns (Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988) and digested by papain to yield Fab and Fc fragments. The Fab and Fc fragments may be separated by affinity chromatography on protein A bead columns.

Monoclonal antibodies of the present invention may be coupled to one or more therapeutic agents. Suitable agents in this regard include radionuclides, differentiation inducers, drugs, toxins, and derivatives thereof. Preferred radionuclides include  $^{90}\text{Y}$ ,  $^{123}\text{I}$ ,  $^{125}\text{I}$ ,  $^{131}\text{I}$ ,  $^{186}\text{Re}$ ,  $^{188}\text{Re}$ ,  $^{211}\text{At}$ , and  $^{212}\text{Bi}$ . Preferred drugs include methotrexate, and pyrimidine and purine analogs. Preferred differentiation inducers include phorbol esters and butyric acid. Preferred toxins include ricin, abrin, diphtheria toxin, cholera toxin, gelonin, *Pseudomonas* exotoxin, *Shigella* toxin, and pokeweed antiviral protein.

A therapeutic agent may be coupled (e.g., covalently bonded) to a suitable monoclonal antibody either directly or indirectly (e.g., via a linker group). A direct reaction between an agent and an antibody is possible when each possesses a substituent capable of reacting with the other. For example, a nucleophilic group, such as an amino or sulfhydryl group, on one may be capable of reacting with a carbonyl-containing group, such as an anhydride or an acid halide, or with an alkyl group containing a good leaving group (e.g., a halide) on the other.

Alternatively, it may be desirable to couple a therapeutic agent and an antibody via a linker group. A linker group can function as a spacer to distance an antibody from an agent in order to avoid interference with binding capabilities. A linker group can also serve to increase the chemical reactivity of a substituent on an agent or an antibody, and



thus increase the coupling efficiency. An increase in chemical reactivity may also facilitate the use of agents, or functional groups on agents, which otherwise would not be possible.

It will be evident to those skilled in the art that a variety of bifunctional or polyfunctional reagents, both homo- and hetero-functional (such as those described in the catalog of the Pierce Chemical Co., Rockford, IL), may be employed as the linker group. Coupling may be effected, for example, through amino groups, carboxyl groups, sulfhydryl groups or oxidized carbohydrate residues. There are numerous references describing such methodology, *e.g.*, U.S. Patent No. 4,671,958, to Rodwell et al.

Where a therapeutic agent is more potent when free from the antibody portion of the immunoconjugates of the present invention, it may be desirable to use a linker group which is cleavable during or upon internalization into a cell. A number of different cleavable linker groups have been described. The mechanisms for the intracellular release of an agent from these linker groups include cleavage by reduction of a disulfide bond (*e.g.*, U.S. Patent No. 4,489,710, to Spitler); by irradiation of a photolabile bond (*e.g.*, U.S. Patent No. 4,625,014, to Senter et al.), by hydrolysis of derivatized amino acid side chains (*e.g.*, U.S. Patent No. 4,638,045, to Kohn et al.), by serum complement-mediated hydrolysis (*e.g.*, U.S. Patent No. 4,671,958, to Rodwell et al.), and acid-catalyzed hydrolysis (*e.g.*, U.S. Patent No. 4,569,789, to Blattler et al.).

It may be desirable to couple more than one agent to an antibody. In one embodiment, multiple molecules of an agent are coupled to one antibody molecule. In another embodiment, more than one type of agent may be coupled to one antibody. Regardless of the particular embodiment, immunoconjugates with more than one agent may be prepared in a variety of ways. For example, more than one agent may be coupled directly to an antibody molecule, or linkers which provide multiple sites for attachment can be used. Alternatively, a carrier can be used.

A carrier may bear the agents in a variety of ways, including covalent bonding either directly or via a linker group. Suitable carriers include proteins such as albumins (*e.g.*, U.S. Patent No. 4,507,234, to Kato et al.), peptides and polysaccharides such as aminodextran (*e.g.*, U.S. Patent No. 4,699,784, to Shih et al.). A carrier may also bear an agent by noncovalent bonding or by encapsulation, such as within a liposome vesicle (*e.g.*, U.S. Patent Nos. 4,429,008 and 4,873,088). Carriers specific for radionuclide agents include radiohalogenated small molecules and chelating compounds. For example, U.S. Patent No. 4,735,792 discloses representative radiohalogenated small molecules and their synthesis. A radionuclide chelate may be formed from chelating compounds that include those containing nitrogen and sulfur atoms as the donor atoms for binding the metal, or metal oxide, radionuclide. For example, U.S. Patent No. 4,673,562, to Davison et al. discloses representative chelating compounds and their synthesis.



A variety of routes of administration for the antibodies and immunoconjugates may be used. Typically, administration will be intravenous, intramuscular, subcutaneous or in the bed of a resected tumor. It will be evident that the precise dose of the antibody/immunoconjugate will vary depending upon the antibody used, the antigen density on the tumor, and the rate of clearance of the antibody.

#### T CELLS

Immunotherapeutic compositions may also, or alternatively, comprise T cells specific for a prostate tumor protein. Such cells may generally be prepared *in vitro* or *ex vivo*, using standard procedures. For example, T cells may be isolated from bone marrow, peripheral blood, or a fraction of bone marrow or peripheral blood of a patient, using a commercially available cell separation system, such as the CEPRATE™ system, available from CellPro Inc., Bothell WA (*see also* U.S. Patent No. 5,240,856; U.S. Patent No. 5,215,926; WO 89/06280; WO 91/16116 and WO 92/07243). Alternatively, T cells may be derived from related or unrelated humans, non-human mammals, cell lines or cultures.

T cells may be stimulated with a prostate tumor polypeptide, polynucleotide encoding a prostate tumor polypeptide and/or an antigen presenting cell (APC) that expresses such a polypeptide. Such stimulation is performed under conditions and for a time sufficient to permit the generation of T cells that are specific for the polypeptide. Preferably, a prostate tumor polypeptide or polynucleotide is present within a delivery vehicle, such as a microsphere, to facilitate the generation of specific T cells.

T cells are considered to be specific for a prostate tumor polypeptide if the T cells kill target cells coated with the polypeptide or expressing a gene encoding the polypeptide. T cell specificity may be evaluated using any of a variety of standard techniques. For example, within a chromium release assay or proliferation assay, a stimulation index of more than two fold increase in lysis and/or proliferation, compared to negative controls, indicates T cell specificity. Such assays may be performed, for example, as described in Chen et al., *Cancer Res.* 54:1065-1070, 1994. Alternatively, detection of the proliferation of T cells may be accomplished by a variety of known techniques. For example, T cell proliferation can be detected by measuring an increased rate of DNA synthesis (*e.g.*, by pulse-labeling cultures of T cells with tritiated thymidine and measuring the amount of tritiated thymidine incorporated into DNA). Contact with a prostate tumor polypeptide (100 ng/ml - 100 µg/ml, preferably 200 ng/ml - 25 µg/ml) for 3 - 7 days should result in at least a two fold increase in proliferation of the T cells. Contact as described above for 2-3 hours should result in activation of the T cells, as measured using standard cytokine assays in which a two fold increase in the level of cytokine release (*e.g.*, TNF or IFN-γ) is indicative of T cell activation (*see* Coligan et al., *Current Protocols in Immunology*, vol. 1, Wiley Interscience

(Greene 1998)). T cells that have been activated in response to a prostate tumor polypeptide, polynucleotide or polypeptide-expressing APC may be CD4<sup>+</sup> and/or CD8<sup>+</sup>. Prostate tumor protein-specific T cells may be expanded using standard techniques. Within preferred embodiments, the T cells are derived from either a patient or a related, or unrelated, donor and are administered to the patient following stimulation and expansion.

For therapeutic purposes, CD4<sup>+</sup> or CD8<sup>+</sup> T cells that proliferate in response to a prostate tumor polypeptide, polynucleotide or APC can be expanded in number either *in vitro* or *in vivo*. Proliferation of such T cells *in vitro* may be accomplished in a variety of ways. For example, the T cells can be re-exposed to a prostate tumor polypeptide, or a short peptide corresponding to an immunogenic portion of such a polypeptide, with or without the addition of T cell growth factors, such as interleukin-2, and/or stimulator cells that synthesize a prostate tumor polypeptide. Alternatively, one or more T cells that proliferate in the presence of a prostate tumor protein can be expanded in number by cloning. Methods for cloning cells are well known in the art, and include limiting dilution.

#### PHARMACEUTICAL COMPOSITIONS AND VACCINES

Within certain aspects, polypeptides, polynucleotides, T cells and/or binding agents disclosed herein may be incorporated into pharmaceutical compositions or immunogenic compositions (*i.e.*, vaccines). Pharmaceutical compositions comprise one or more such compounds and a physiologically acceptable carrier. Vaccines may comprise one or more such compounds and a non-specific immune response enhancer. A non-specific immune response enhancer may be any substance that enhances an immune response to an exogenous antigen. Examples of non-specific immune response enhancers include adjuvants, biodegradable microspheres (*e.g.*, polylactic galactide) and liposomes (into which the compound is incorporated; *see e.g.*, Fullerton, U.S. Patent No. 4,235,877). Vaccine preparation is generally described in, for example, M.F. Powell and M.J. Newman, eds., "Vaccine Design (the subunit and adjuvant approach)," Plenum Press (NY, 1995). Pharmaceutical compositions and vaccines within the scope of the present invention may also contain other compounds, which may be biologically active or inactive. For example, one or more immunogenic portions of other tumor antigens may be present, either incorporated into a fusion polypeptide or as a separate compound, within the composition or vaccine.

A pharmaceutical composition or vaccine may contain DNA encoding one or more of the polypeptides as described above, such that the polypeptide is generated *in situ*. As noted above, the DNA may be present within any of a variety of delivery systems known to those of ordinary skill in the art, including nucleic acid expression systems, bacteria and viral expression systems. Numerous gene delivery techniques are well known in the art, such as those described by Rolland, *Crit. Rev. Therap. Drug Carrier Systems* 15:143-198, 1998,

and references cited therein. Appropriate nucleic acid expression systems contain the necessary DNA sequences for expression in the patient (such as a suitable promoter and terminating signal). Bacterial delivery systems involve the administration of a bacterium (such as *Bacillus-Calmette-Guerrin*) that expresses an immunogenic portion of the polypeptide on its cell surface or secretes such an epitope. In a preferred embodiment, the DNA may be introduced using a viral expression system (e.g., vaccinia or other pox virus, retrovirus, or adenovirus), which may involve the use of a non-pathogenic (defective), replication competent virus. Suitable systems are disclosed, for example, in Fisher-Hoch et al., *Proc. Natl. Acad. Sci. USA* 86:317-321, 1989; Flexner et al., *Ann. N.Y. Acad. Sci.* 569:86-103, 1989; Flexner et al., *Vaccine* 8:17-21, 1990; U.S. Patent Nos. 4,603,112, 4,769,330, and 5,017,487; WO 89/01973; U.S. Patent No. 4,777,127; GB 2,200,651; EP 0,345,242; WO 91/02805; Berkner, *Biotechniques* 6:616-627, 1988; Rosenfeld et al., *Science* 252:431-434, 1991; Kolls et al., *Proc. Natl. Acad. Sci. USA* 91:215-219, 1994; Kass-Eisler et al., *Proc. Natl. Acad. Sci. USA* 90:11498-11502, 1993; Guzman et al., *Circulation* 88:2838-2848, 1993; and Guzman et al., *Cir. Res.* 73:1202-1207, 1993. Techniques for incorporating DNA into such expression systems are well known to those of ordinary skill in the art. The DNA may also be "naked," as described, for example, in Ulmer et al., *Science* 259:1745-1749, 1993 and reviewed by Cohen, *Science* 259:1691-1692, 1993. The uptake of naked DNA may be increased by coating the DNA onto biodegradable beads, which are efficiently transported into the cells.

While any suitable carrier known to those of ordinary skill in the art may be employed in the pharmaceutical compositions of this invention, the type of carrier will vary depending on the mode of administration. Compositions of the present invention may be formulated for any appropriate manner of administration, including for example, topical, oral, nasal, intravenous, intracranial, intraperitoneal, subcutaneous or intramuscular administration. For parenteral administration, such as subcutaneous injection, the carrier preferably comprises water, saline, alcohol, a fat, a wax or a buffer. For oral administration, any of the above carriers or a solid carrier, such as mannitol, lactose, starch, magnesium stearate, sodium saccharine, talcum, cellulose, glucose, sucrose, and magnesium carbonate, may be employed. Biodegradable microspheres (e.g., polylactate polyglycolate) may also be employed as carriers for the pharmaceutical compositions of this invention. Suitable biodegradable microspheres are disclosed, for example, in U.S. Patent Nos. 4,897,268 and 5,075,109.

Such compositions may also comprise buffers (e.g., neutral buffered saline or phosphate buffered saline), carbohydrates (e.g., glucose, mannose, sucrose or dextrans), mannitol, proteins, polypeptides or amino acids such as glycine, antioxidants, chelating agents such as EDTA or glutathione, adjuvants (e.g., aluminum hydroxide) and/or

preservatives. Alternatively, compositions of the present invention may be formulated as a lyophilizate. Compounds may also be encapsulated within liposomes using well known technology.

Any of a variety of non-specific immune response enhancers may be employed in the vaccines of this invention. For example, an adjuvant may be included. Most adjuvants contain a substance designed to protect the antigen from rapid catabolism, such as aluminum hydroxide or mineral oil, and a stimulator of immune responses, such as lipid A, *Bordetella pertussis* or *Mycobacterium tuberculosis* derived proteins. Suitable adjuvants are commercially available as, for example, Freund's Incomplete Adjuvant and Complete Adjuvant (Difco Laboratories, Detroit, MI); Merck Adjuvant 65 (Merck and Company, Inc., Rahway, NJ); aluminum salts such as aluminum hydroxide gel (alum) or aluminum phosphate; salts of calcium, iron or zinc; an insoluble suspension of acylated tyrosine; acylated sugars; cationically or anionically derivatized polysaccharides; polyphosphazenes; biodegradable microspheres; monophosphoryl lipid A and quil A. Cytokines, such as GM-CSF or interleukin-2, -7, or -12, may also be used as adjuvants.

Within the vaccines provided herein, the adjuvant composition is preferably designed to induce an immune response predominantly of the Th1 type. High levels of Th1-type cytokines (e.g., IFN- $\gamma$ , IL-2 and IL-12) tend to favor the induction of cell mediated immune responses to an administered antigen. In contrast, high levels of Th2-type cytokines (e.g., IL-4, IL-5, IL-6, IL-10 and TNF- $\beta$ ) tend to favor the induction of humoral immune responses. Following application of a vaccine as provided herein, a patient will support an immune response that includes Th1- and Th2-type responses. Within a preferred embodiment, in which a response is predominantly Th1-type, the level of Th1-type cytokines will increase to a greater extent than the level of Th2-type cytokines. The levels of these cytokines may be readily assessed using standard assays. For a review of the families of cytokines, see Mosmann and Coffman, *Ann. Rev. Immunol.* 7:145-173, 1989.

Preferred adjuvants for use in eliciting a predominantly Th1-type response include, for example, a combination of monophosphoryl lipid A, preferably 3-de-O-acylated monophosphoryl lipid A (3D-MPL), together with an aluminum salt. MPL adjuvants are available from Ribi ImmunoChem Research Inc. (Hamilton, MT; see US Patent Nos. 4,436,727; 4,877,611; 4,866,034 and 4,912,094). CpG-containing oligonucleotides (in which the CpG dinucleotide is unmethylated) also induce a predominantly Th1 response. Such oligonucleotides are well known and are described, for example, in WO 96/02555. Another preferred adjuvant is a saponin, preferably QS21, which may be used alone or in combination with other adjuvants. For example, an enhanced system involves the combination of a monophosphoryl lipid A and saponin derivative, such as the combination of QS21 and 3D-MPL as described in WO 94/00153, or a less reactogenic composition where the QS21 is

quenched with cholesterol, as described in WO 96/33739. Other preferred formulations comprises an oil-in-water emulsion and tocopherol. A particularly potent adjuvant formulation involving QS21, 3D-MPL and tocopherol in an oil-in-water emulsion is described in WO 95/17210. Any vaccine provided herein may be prepared using well known methods that result in a combination of antigen, immune response enhancer and a suitable carrier or excipient.

The compositions described herein may be administered as part of a sustained release formulation (*i.e.*, a formulation such as a capsule or sponge that effects a slow release of compound following administration). Such formulations may generally be prepared using well known technology and administered by, for example, oral, rectal or subcutaneous implantation, or by implantation at the desired target site. Sustained-release formulations may contain a polypeptide, polynucleotide or antibody dispersed in a carrier matrix and/or contained within a reservoir surrounded by a rate controlling membrane. Carriers for use within such formulations are biocompatible, and may also be biodegradable; preferably the formulation provides a relatively constant level of active component release. The amount of active compound contained within a sustained release formulation depends upon the site of implantation, the rate and expected duration of release and the nature of the condition to be treated or prevented.

Any of a variety of delivery vehicles may be employed within pharmaceutical compositions and vaccines to facilitate production of an antigen-specific immune response that targets tumor cells. Delivery vehicles include antigen presenting cells (APCs), such as dendritic cells, macrophages, B cells, monocytes and other cells that may be engineered to be efficient APCs. Such cells may, but need not, be genetically modified to increase the capacity for presenting the antigen, to improve activation and/or maintenance of the T cell response, to have anti-tumor effects *per se* and/or to be immunologically compatible with the receiver (*i.e.*, matched HLA haplotype). APCs may generally be isolated from any of a variety of biological fluids and organs, including tumor and peritumoral tissues, and may be autologous, allogeneic, syngeneic or xenogeneic cells.

Certain preferred embodiments of the present invention use dendritic cells or progenitors thereof as antigen-presenting cells. Dendritic cells are highly potent APCs (Banchereau and Steinman, *Nature* 392:245-251, 1998) and have been shown to be effective as a physiological adjuvant for eliciting prophylactic or therapeutic antitumor immunity (*see* Timmerman and Levy, *Ann. Rev. Med.* 50:507-529, 1999). In general, dendritic cells may be identified based on their typical shape (stellate *in situ*, with marked cytoplasmic processes (dendrites) visible *in vitro*) and based on the lack of differentiation markers of B cells (CD19 and CD20), T cells (CD3), monocytes (CD14) and natural killer cells (CD56), as determined using standard assays. Dendritic cells may, of course, be engineered to express specific cell-

surface receptors or ligands that are not commonly found on dendritic cells *in vivo* or *ex vivo*, and such modified dendritic cells are contemplated by the present invention. As an alternative to dendritic cells, secreted vesicles antigen-loaded dendritic cells (called exosomes) may be used within a vaccine (see Zitvogel et al., *Nature Med.* 4:594-600, 1998).

Dendritic cells and progenitors may be obtained from peripheral blood, bone marrow, tumor-infiltrating cells, peritumoral tissues-infiltrating cells, lymph nodes, spleen, skin, umbilical cord blood or any other suitable tissue or fluid. For example, dendritic cells may be differentiated *ex vivo* by adding a combination of cytokines such as GM-CSF, IL-4, IL-13 and/or TNF $\alpha$  to cultures of monocytes harvested from peripheral blood. Alternatively, CD34 positive cells harvested from peripheral blood, umbilical cord blood or bone marrow may be differentiated into dendritic cells by adding to the culture medium combinations of GM-CSF, IL-3, TNF $\alpha$ , CD40 ligand, LPS, flt3 ligand and/or other compound(s) that induce maturation and proliferation of dendritic cells.

Dendritic cells are conveniently categorized as "immature" and "mature" cells, which allows a simple way to discriminate between two well characterized phenotypes. However, this nomenclature should not be construed to exclude all possible intermediate stages of differentiation. Immature dendritic cells are characterized as APC with a high capacity for antigen uptake and processing, which correlates with the high expression of Fc $\gamma$  receptor, mannose receptor and DEC-205 marker. The mature phenotype is typically characterized by a lower expression of these markers, but a high expression of cell surface molecules responsible for T cell activation such as class I and class II MHC, adhesion molecules (e.g., CD54 and CD11) and costimulatory molecules (e.g., CD40, CD80 and CD86).

APCs may generally be transfected with a polynucleotide encoding a prostate tumor protein (or portion or other variant thereof) such that the prostate tumor polypeptide, or an immunogenic portion thereof, is expressed on the cell surface. Such transfection may take place *ex vivo*, and a composition or vaccine comprising such transfected cells may then be used for therapeutic purposes, as described herein. Alternatively, a gene delivery vehicle that targets a dendritic or other antigen presenting cell may be administered to a patient, resulting in transfection that occurs *in vivo*. *In vivo* and *ex vivo* transfection of dendritic cells, for example, may generally be performed using any methods known in the art, such as those described in WO 97/24447, or the gene gun approach described by Mahvi et al., *Immunology and cell Biology* 75:456-460, 1997. Antigen loading of dendritic cells may be achieved by incubating dendritic cells or progenitor cells with the prostate tumor polypeptide, DNA (naked or within a plasmid vector) or RNA; or with antigen-expressing recombinant bacterium or viruses (e.g., vaccinia, fowlpox, adenovirus or lentivirus vectors). Prior to loading, the polypeptide may be covalently conjugated to an immunological partner that

provides T cell help (e.g., a carrier molecule). Alternatively, a dendritic cell may be pulsed with a non-conjugated immunological partner, separately or in the presence of the polypeptide.

#### CANCER THERAPY

In further aspects of the present invention, the compositions described herein may be used for immunotherapy of cancer, such as prostate cancer. Within such methods, pharmaceutical compositions and vaccines are typically administered to a patient. As used herein, a "patient" refers to any warm-blooded animal, preferably a human. A patient may or may not be afflicted with cancer. Accordingly, the above pharmaceutical compositions and vaccines may be used to prevent the development of a cancer or to treat a patient afflicted with a cancer. A cancer may be diagnosed using criteria generally accepted in the art, including the presence of a malignant tumor. Pharmaceutical compositions and vaccines may be administered either prior to or following surgical removal of primary tumors and/or treatment such as administration of radiotherapy or conventional chemotherapeutic drugs.

Within certain embodiments, immunotherapy may be active immunotherapy, in which treatment relies on the *in vivo* stimulation of the endogenous host immune system to react against tumors with the administration of immune response-modifying agents (such as polypeptides and polynucleotides disclosed herein).

Within other embodiments, immunotherapy may be passive immunotherapy, in which treatment involves the delivery of agents with established tumor-immune reactivity (such as effector cells or antibodies) that can directly or indirectly mediate antitumor effects and does not necessarily depend on an intact host immune system. Examples of effector cells include T cells as discussed above, T lymphocytes (such as CD8<sup>+</sup> cytotoxic T lymphocytes and CD4<sup>+</sup> T-helper tumor-infiltrating lymphocytes), killer cells (such as Natural Killer cells and lymphokine-activated killer cells), B cells and antigen-presenting cells (such as dendritic cells and macrophages) expressing a polypeptide provided herein. T cell receptors and antibody receptors specific for the polypeptides recited herein may be cloned, expressed and transferred into other vectors or effector cells for adoptive immunotherapy. The polypeptides provided herein may also be used to generate antibodies or anti-idiotypic antibodies (as described above and in U.S. Patent No. 4,918,164) for passive immunotherapy.

Effector cells may generally be obtained in sufficient quantities for adoptive immunotherapy by growth *in vitro*, as described herein. Culture conditions for expanding single antigen-specific effector cells to several billion in number with retention of antigen recognition *in vivo* are well known in the art. Such *in vitro* culture conditions typically use intermittent stimulation with antigen, often in the presence of cytokines (such as IL-2) and non-dividing feeder cells. As noted above, immunoreactive polypeptides as provided herein

may be used to rapidly expand antigen-specific T cell cultures in order to generate a sufficient number of cells for immunotherapy. In particular, antigen-presenting cells, such as dendritic, macrophage, monocyte, fibroblast or B cells, may be pulsed with immunoreactive polypeptides or transfected with one or more polynucleotides using standard techniques well known in the art. For example, antigen-presenting cells can be transfected with a polynucleotide having a promoter appropriate for increasing expression in a recombinant virus or other expression system. Cultured effector cells for use in therapy must be able to grow and distribute widely, and to survive long term *in vivo*. Studies have shown that cultured effector cells can be induced to grow *in vivo* and to survive long term in substantial numbers by repeated stimulation with antigen supplemented with IL-2 (*see, for example, Cheever et al., Immunological Reviews 157:177, 1997*).

Alternatively, a vector expressing a polypeptide recited herein may be introduced into antigen presenting cells taken from a patient and clonally propagated *ex vivo* for transplant back into the same patient. Transfected cells may be reintroduced into the patient using any means known in the art, preferably in sterile form by intravenous, intracavitary, intraperitoneal or intratumor administration.

Routes and frequency of administration of the therapeutic compositions disclosed herein, as well as dosage, will vary from individual to individual, and may be readily established using standard techniques. In general, the pharmaceutical compositions and vaccines may be administered by injection (*e.g.*, intracutaneous, intramuscular, intravenous or subcutaneous), intranasally (*e.g.*, by aspiration) or orally. Preferably, between 1 and 10 doses may be administered over a 52 week period. Preferably, 6 doses are administered, at intervals of 1 month, and booster vaccinations may be given periodically thereafter. Alternate protocols may be appropriate for individual patients. A suitable dose is an amount of a compound that, when administered as described above, is capable of promoting an anti-tumor immune response, and is at least 10-50% above the basal (*i.e.*, untreated) level. Such response can be monitored by measuring the anti-tumor antibodies in a patient or by vaccine-dependent generation of cytolytic effector cells capable of killing the patient's tumor cells *in vitro*. Such vaccines should also be capable of causing an immune response that leads to an improved clinical outcome (*e.g.*, more frequent remissions, complete or partial or longer disease-free survival) in vaccinated patients as compared to non-vaccinated patients. In general, for pharmaceutical compositions and vaccines comprising one or more polypeptides, the amount of each polypeptide present in a dose ranges from about 100  $\mu$ g to 5 mg per kg of host. Suitable dose sizes will vary with the size of the patient, but will typically range from about 0.1 mL to about 5 mL.

In general, an appropriate dosage and treatment regimen provides the active compound(s) in an amount sufficient to provide therapeutic and/or prophylactic benefit. Such



a response can be monitored by establishing an improved clinical outcome (e.g., more frequent remissions, complete or partial, or longer disease-free survival) in treated patients as compared to non-treated patients. Increases in preexisting immune responses to a prostate tumor protein generally correlate with an improved clinical outcome. Such immune responses may generally be evaluated using standard proliferation, cytotoxicity or cytokine assays, which may be performed using samples obtained from a patient before and after treatment.

#### METHODS FOR DETECTING CANCER

In general, a cancer may be detected in a patient based on the presence of one or more prostate tumor proteins and/or polynucleotides encoding such proteins in a biological sample (for example, blood, sera, urine and/or tumor biopsies) obtained from the patient. In other words, such proteins may be used as markers to indicate the presence or absence of a cancer such as prostate cancer. In addition, such proteins may be useful for the detection of other cancers. The binding agents provided herein generally permit detection of the level of antigen that binds to the agent in the biological sample. Polynucleotide primers and probes may be used to detect the level of mRNA encoding a tumor protein, which is also indicative of the presence or absence of a cancer. In general, a prostate tumor sequence should be present at a level that is at least three fold higher in tumor tissue than in normal tissue.

There are a variety of assay formats known to those of ordinary skill in the art for using a binding agent to detect polypeptide markers in a sample. See, e.g., Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. In general, the presence or absence of a cancer in a patient may be determined by (a) contacting a biological sample obtained from a patient with a binding agent; (b) detecting in the sample a level of polypeptide that binds to the binding agent; and (c) comparing the level of polypeptide with a predetermined cut-off value.

In a preferred embodiment, the assay involves the use of binding agent immobilized on a solid support to bind to and remove the polypeptide from the remainder of the sample. The bound polypeptide may then be detected using a detection reagent that contains a reporter group and specifically binds to the binding agent/polypeptide complex. Such detection reagents may comprise, for example, a binding agent that specifically binds to the polypeptide or an antibody or other agent that specifically binds to the binding agent, such as an anti-immunoglobulin, protein G, protein A or a lectin. Alternatively, a competitive assay may be utilized, in which a polypeptide is labeled with a reporter group and allowed to bind to the immobilized binding agent after incubation of the binding agent with the sample. The extent to which components of the sample inhibit the binding of the labeled polypeptide to the binding agent is indicative of the reactivity of the sample with the immobilized binding

agent. Suitable polypeptides for use within such assays include full length prostate tumor proteins and portions thereof to which the binding agent binds, as described above.

The solid support may be any material known to those of ordinary skill in the art to which the tumor protein may be attached. For example, the solid support may be a test well in a microtiter plate or a nitrocellulose or other suitable membrane. Alternatively, the support may be a bead or disc, such as glass, fiberglass, latex or a plastic material such as polystyrene or polyvinylchloride. The support may also be a magnetic particle or a fiber optic sensor, such as those disclosed, for example, in U.S. Patent No. 5,359,681. The binding agent may be immobilized on the solid support using a variety of techniques known to those of skill in the art, which are amply described in the patent and scientific literature. In the context of the present invention, the term "immobilization" refers to both noncovalent association, such as adsorption, and covalent attachment (which may be a direct linkage between the agent and functional groups on the support or may be a linkage by way of a cross-linking agent). Immobilization by adsorption to a well in a microtiter plate or to a membrane is preferred. In such cases, adsorption may be achieved by contacting the binding agent, in a suitable buffer, with the solid support for a suitable amount of time. The contact time varies with temperature, but is typically between about 1 hour and about 1 day. In general, contacting a well of a plastic microtiter plate (such as polystyrene or polyvinylchloride) with an amount of binding agent ranging from about 10 ng to about 10  $\mu$ g, and preferably about 100 ng to about 1  $\mu$ g, is sufficient to immobilize an adequate amount of binding agent.

Covalent attachment of binding agent to a solid support may generally be achieved by first reacting the support with a bifunctional reagent that will react with both the support and a functional group, such as a hydroxyl or amino group, on the binding agent. For example, the binding agent may be covalently attached to supports having an appropriate polymer coating using benzoquinone or by condensation of an aldehyde group on the support with an amine and an active hydrogen on the binding partner (*see, e.g.,* Pierce Immunotechnology Catalog and Handbook, 1991, at A12-A13).

In certain embodiments, the assay is a two-antibody sandwich assay. This assay may be performed by first contacting an antibody that has been immobilized on a solid support, commonly the well of a microtiter plate, with the sample, such that polypeptides within the sample are allowed to bind to the immobilized antibody. Unbound sample is then removed from the immobilized polypeptide-antibody complexes and a detection reagent (preferably a second antibody capable of binding to a different site on the polypeptide) containing a reporter group is added. The amount of detection reagent that remains bound to the solid support is then determined using a method appropriate for the specific reporter group.

More specifically, once the antibody is immobilized on the support as described above, the remaining protein binding sites on the support are typically blocked. Any suitable blocking agent known to those of ordinary skill in the art, such as bovine serum albumin or Tween 20™ (Sigma Chemical Co., St. Louis, MO). The immobilized antibody is then incubated with the sample, and polypeptide is allowed to bind to the antibody. The sample may be diluted with a suitable diluent, such as phosphate-buffered saline (PBS) prior to incubation. In general, an appropriate contact time (*i.e.*, incubation time) is a period of time that is sufficient to detect the presence of polypeptide within a sample obtained from an individual with prostate cancer. Preferably, the contact time is sufficient to achieve a level of binding that is at least about 95% of that achieved at equilibrium between bound and unbound polypeptide. Those of ordinary skill in the art will recognize that the time necessary to achieve equilibrium may be readily determined by assaying the level of binding that occurs over a period of time. At room temperature, an incubation time of about 30 minutes is generally sufficient.

Unbound sample may then be removed by washing the solid support with an appropriate buffer, such as PBS containing 0.1% Tween 20™. The second antibody, which contains a reporter group, may then be added to the solid support. Preferred reporter groups include those groups recited above.

The detection reagent is then incubated with the immobilized antibody-polypeptide complex for an amount of time sufficient to detect the bound polypeptide. An appropriate amount of time may generally be determined by assaying the level of binding that occurs over a period of time. Unbound detection reagent is then removed and bound detection reagent is detected using the reporter group. The method employed for detecting the reporter group depends upon the nature of the reporter group. For radioactive groups, scintillation counting or autoradiographic methods are generally appropriate. Spectroscopic methods may be used to detect dyes, luminescent groups and fluorescent groups. Biotin may be detected using avidin, coupled to a different reporter group (commonly a radioactive or fluorescent group or an enzyme). Enzyme reporter groups may generally be detected by the addition of substrate (generally for a specific period of time), followed by spectroscopic or other analysis of the reaction products.

To determine the presence or absence of a cancer, such as prostate cancer, the signal detected from the reporter group that remains bound to the solid support is generally compared to a signal that corresponds to a predetermined cut-off value. In one preferred embodiment, the cut-off value for the detection of a cancer is the average mean signal obtained when the immobilized antibody is incubated with samples from patients without the cancer. In general, a sample generating a signal that is three standard deviations above the predetermined cut-off value is considered positive for the cancer. In an alternate preferred

embodiment, the cut-off value is determined using a Receiver Operator Curve, according to the method of Sackett et al., *Clinical Epidemiology: A Basic Science for Clinical Medicine*, Little Brown and Co., 1985, p. 106-7. Briefly, in this embodiment, the cut-off value may be determined from a plot of pairs of true positive rates (*i.e.*, sensitivity) and false positive rates (100%-specificity) that correspond to each possible cut-off value for the diagnostic test result. The cut-off value on the plot that is the closest to the upper left-hand corner (*i.e.*, the value that encloses the largest area) is the most accurate cut-off value, and a sample generating a signal that is higher than the cut-off value determined by this method may be considered positive. Alternatively, the cut-off value may be shifted to the left along the plot, to minimize the false positive rate, or to the right, to minimize the false negative rate. In general, a sample generating a signal that is higher than the cut-off value determined by this method is considered positive for a cancer.

In a related embodiment, the assay is performed in a flow-through or strip test format, wherein the binding agent is immobilized on a membrane, such as nitrocellulose. In the flow-through test, polypeptides within the sample bind to the immobilized binding agent as the sample passes through the membrane. A second, labeled binding agent then binds to the binding agent-polypeptide complex as a solution containing the second binding agent flows through the membrane. The detection of bound second binding agent may then be performed as described above. In the strip test format, one end of the membrane to which binding agent is bound is immersed in a solution containing the sample. The sample migrates along the membrane through a region containing second binding agent and to the area of immobilized binding agent. Concentration of second binding agent at the area of immobilized antibody indicates the presence of a cancer. Typically, the concentration of second binding agent at that site generates a pattern, such as a line, that can be read visually. The absence of such a pattern indicates a negative result. In general, the amount of binding agent immobilized on the membrane is selected to generate a visually discernible pattern when the biological sample contains a level of polypeptide that would be sufficient to generate a positive signal in the two-antibody sandwich assay, in the format discussed above. Preferred binding agents for use in such assays are antibodies and antigen-binding fragments thereof. Preferably, the amount of antibody immobilized on the membrane ranges from about 25 ng to about 1  $\mu$ g, and more preferably from about 50 ng to about 500 ng. Such tests can typically be performed with a very small amount of biological sample.

Of course, numerous other assay protocols exist that are suitable for use with the tumor proteins or binding agents of the present invention. The above descriptions are intended to be exemplary only. For example, it will be apparent to those of ordinary skill in the art that the above protocols may be readily modified to use prostate tumor polypeptides to

detect antibodies that bind to such polypeptides in a biological sample. The detection of such prostate tumor protein specific antibodies may correlate with the presence of a cancer.

A cancer may also, or alternatively, be detected based on the presence of T cells that specifically react with a prostate tumor protein in a biological sample. Within certain methods, a biological sample comprising CD4<sup>+</sup> and/or CD8<sup>+</sup> T cells isolated from a patient is incubated with a prostate tumor polypeptide, a polynucleotide encoding such a polypeptide and/or an APC that expresses at least an immunogenic portion of such a polypeptide, and the presence or absence of specific activation of the T cells is detected. Suitable biological samples include, but are not limited to, isolated T cells. For example, T cells may be isolated from a patient by routine techniques (such as by Ficoll/Hypaque density gradient centrifugation of peripheral blood lymphocytes). T cells may be incubated *in vitro* for 2-9 days (typically 4 days) at 37°C with prostate tumor polypeptide (*e.g.*, 5 - 25 µg/ml). It may be desirable to incubate another aliquot of a T cell sample in the absence of prostate tumor polypeptide to serve as a control. For CD4<sup>+</sup> T cells, activation is preferably detected by evaluating proliferation of the T cells. For CD8<sup>+</sup> T cells, activation is preferably detected by evaluating cytolytic activity. A level of proliferation that is at least two fold greater and/or a level of cytolytic activity that is at least 20% greater than in disease-free patients indicates the presence of a cancer in the patient.

As noted above, a cancer may also, or alternatively, be detected based on the level of mRNA encoding a prostate tumor protein in a biological sample. For example, at least two oligonucleotide primers may be employed in a polymerase chain reaction (PCR) based assay to amplify a portion of a prostate tumor cDNA derived from a biological sample, wherein at least one of the oligonucleotide primers is specific for (*i.e.*, hybridizes to) a polynucleotide encoding the prostate tumor protein. The amplified cDNA is then separated and detected using techniques well known in the art, such as gel electrophoresis. Similarly, oligonucleotide probes that specifically hybridize to a polynucleotide encoding a prostate tumor protein may be used in a hybridization assay to detect the presence of polynucleotide encoding the tumor protein in a biological sample.

To permit hybridization under assay conditions, oligonucleotide primers and probes should comprise an oligonucleotide sequence that has at least about 60%, preferably at least about 75% and more preferably at least about 90%, identity to a portion of a polynucleotide encoding a prostate tumor protein that is at least 10 nucleotides, and preferably at least 20 nucleotides, in length. Preferably, oligonucleotide primers and/or probes will hybridize to a polynucleotide encoding a polypeptide disclosed herein under moderately stringent conditions, as defined above. Oligonucleotide primers and/or probes which may be usefully employed in the diagnostic methods described herein preferably are at least 10-40 nucleotides in length. In a preferred embodiment, the oligonucleotide primers

comprise at least 10 contiguous nucleotides, more preferably at least 15 contiguous nucleotides, of a DNA molecule having a sequence recited in SEQ ID NO: 1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375 and 381. Techniques for both PCR based assays and hybridization assays are well known in the art (*see, for example, Mullis et al., Cold Spring Harbor Symp. Quant. Biol., 51:263, 1987; Erlich ed., PCR Technology, Stockton Press, NY, 1989*).

One preferred assay employs RT-PCR, in which PCR is applied in conjunction with reverse transcription. Typically, RNA is extracted from a biological sample, such as biopsy tissue, and is reverse transcribed to produce cDNA molecules. PCR amplification using at least one specific primer generates a cDNA molecule, which may be separated and visualized using, for example, gel electrophoresis. Amplification may be performed on biological samples taken from a test patient and from an individual who is not afflicted with a cancer. The amplification reaction may be performed on several dilutions of cDNA spanning two orders of magnitude. A two-fold or greater increase in expression in several dilutions of the test patient sample as compared to the same dilutions of the non-cancerous sample is typically considered positive.

In another embodiment, the disclosed compositions may be used as markers for the progression of cancer. In this embodiment, assays as described above for the diagnosis of a cancer may be performed over time, and the change in the level of reactive polypeptide(s) or polynucleotide evaluated. For example, the assays may be performed every 24-72 hours for a period of 6 months to 1 year, and thereafter performed as needed. In general, a cancer is progressing in those patients in whom the level of polypeptide or polynucleotide detected increases over time. In contrast, the cancer is not progressing when the level of reactive polypeptide or polynucleotide either remains constant or decreases with time.

Certain *in vivo* diagnostic assays may be performed directly on a tumor. One such assay involves contacting tumor cells with a binding agent. The bound binding agent may then be detected directly or indirectly via a reporter group. Such binding agents may also be used in histological applications. Alternatively, polynucleotide probes may be used within such applications.

As noted above, to improve sensitivity, multiple prostate tumor protein markers may be assayed within a given sample. It will be apparent that binding agents specific for different proteins provided herein may be combined within a single assay. Further, multiple primers or probes may be used concurrently. The selection of tumor protein markers may be based on routine experiments to determine combinations that results in optimal sensitivity. In addition, or alternatively, assays for tumor proteins provided herein may be combined with assays for other known tumor antigens.

## DIAGNOSTIC KITS

The present invention further provides kits for use within any of the above diagnostic methods. Such kits typically comprise two or more components necessary for performing a diagnostic assay. Components may be compounds, reagents, containers and/or equipment. For example, one container within a kit may contain a monoclonal antibody or fragment thereof that specifically binds to a prostate tumor protein. Such antibodies or fragments may be provided attached to a support material, as described above. One or more additional containers may enclose elements, such as reagents or buffers, to be used in the assay. Such kits may also, or alternatively, contain a detection reagent as described above that contains a reporter group suitable for direct or indirect detection of antibody binding.

Alternatively, a kit may be designed to detect the level of mRNA encoding a prostate tumor protein in a biological sample. Such kits generally comprise at least one oligonucleotide probe or primer, as described above, that hybridizes to a polynucleotide encoding a prostate tumor protein. Such an oligonucleotide may be used, for example, within a PCR or hybridization assay. Additional components that may be present within such kits include a second oligonucleotide and/or a diagnostic reagent or container to facilitate the detection of a polynucleotide encoding a prostate tumor protein.

The following Examples are offered by way of illustration and not by way of limitation.

## EXAMPLES

### EXAMPLE 1

#### ISOLATION AND CHARACTERIZATION OF PROSTATE TUMOR POLYPEPTIDES

This Example describes the isolation of certain prostate tumor polypeptides from a prostate tumor cDNA library.

A human prostate tumor cDNA expression library was constructed from prostate tumor poly A<sup>+</sup> RNA using a Superscript Plasmid System for cDNA Synthesis and Plasmid Cloning kit (BRL Life Technologies, Gaithersburg, MD 20897) following the manufacturer's protocol. Specifically, prostate tumor tissues were homogenized with polytron (Kinematica, Switzerland) and total RNA was extracted using Trizol reagent (BRL Life Technologies) as directed by the manufacturer. The poly A<sup>+</sup> RNA was then purified using a Qiagen oligotex spin column mRNA purification kit (Qiagen, Santa Clarita, CA 91355) according to the manufacturer's protocol. First-strand cDNA was synthesized using the NotI/Oligo-dT18 primer. Double-stranded cDNA was synthesized, ligated with EcoRI/BAXI adaptors (Invitrogen, San Diego, CA) and digested with NotI. Following size fractionation with Chroma Spin-1000 columns (Clontech, Palo Alto, CA), the cDNA was ligated into the EcoRI/NotI site of pCDNA3.1 (Invitrogen) and transformed into ElectroMax *E. coli* DH10B cells (BRL Life Technologies) by electroporation.

Using the same procedure, a normal human pancreas cDNA expression library was prepared from a pool of six tissue specimens (Clontech). The cDNA libraries were characterized by determining the number of independent colonies, the percentage of clones that carried insert, the average insert size and by sequence analysis. The prostate tumor library contained  $1.64 \times 10^7$  independent colonies, with 70% of clones having an insert and the average insert size being 1745 base pairs. The normal pancreas cDNA library contained  $3.3 \times 10^6$  independent colonies, with 69% of clones having inserts and the average insert size being 1120 base pairs. For both libraries, sequence analysis showed that the majority of clones had a full length cDNA sequence and were synthesized from mRNA, with minimal rRNA and mitochondrial DNA contamination.

cDNA library subtraction was performed using the above prostate tumor and normal pancreas cDNA libraries, as described by Hara *et al.* (*Blood*, 84:189-199, 1994) with some modifications. Specifically, a prostate tumor-specific subtracted cDNA library was generated as follows. Normal pancreas cDNA library (70 µg) was digested with EcoRI, NotI, and SfuI, followed by a filling-in reaction with DNA polymerase Klenow fragment. After phenol-chloroform extraction and ethanol precipitation, the DNA was dissolved in 100 µl of



H<sub>2</sub>O, heat-denatured and mixed with 100 µl (100 µg) of Photoprobe biotin (Vector Laboratories, Burlingame, CA). As recommended by the manufacturer, the resulting mixture was irradiated with a 270 W sunlamp on ice for 20 minutes. Additional Photoprobe biotin (50 µl) was added and the biotinylation reaction was repeated. After extraction with butanol five times, the DNA was ethanol-precipitated and dissolved in 23 µl H<sub>2</sub>O to form the driver DNA.

To form the tracer DNA, 10 µg prostate tumor cDNA library was digested with BamHI and XhoI, phenol chloroform extracted and passed through Chroma spin-400 columns (Clontech). Following ethanol precipitation, the tracer DNA was dissolved in 5 µl H<sub>2</sub>O. Tracer DNA was mixed with 15 µl driver DNA and 20 µl of 2 x hybridization buffer (1.5 M NaCl/10 mM EDTA/50 mM HEPES pH 7.5/0.2% sodium dodecyl sulfate), overlaid with mineral oil, and heat-denatured completely. The sample was immediately transferred into a 68 °C water bath and incubated for 20 hours (long hybridization [LH]). The reaction mixture was then subjected to a streptavidin treatment followed by phenol/chloroform extraction. This process was repeated three more times. Subtracted DNA was precipitated, dissolved in 12 µl H<sub>2</sub>O, mixed with 8 µl driver DNA and 20 µl of 2 x hybridization buffer, and subjected to a hybridization at 68 °C for 2 hours (short hybridization [SH]). After removal of biotinylated double-stranded DNA, subtracted cDNA was ligated into BamHI/XhoI site of chloramphenicol resistant pBCSK<sup>+</sup> (Stratagene, La Jolla, CA 92037) and transformed into ElectroMax *E. coli* DH10B cells by electroporation to generate a prostate tumor specific subtracted cDNA library (referred to as "prostate subtraction 1").

To analyze the subtracted cDNA library, plasmid DNA was prepared from 100 independent clones, randomly picked from the subtracted prostate tumor specific library and grouped based on insert size. Representative cDNA clones were further characterized by DNA sequencing with a Perkin Elmer/Applied Biosystems Division Automated Sequencer Model 373A (Foster City, CA). Six cDNA clones, hereinafter referred to as F1-13, F1-12, F1-16, H1-1, H1-9 and H1-4, were shown to be abundant in the subtracted prostate-specific cDNA library. The determined 3' and 5' cDNA sequences for F1-12 are provided in SEQ ID NO: 2 and 3, respectively, with determined 3' cDNA sequences for F1-13, F1-16, H1-1, H1-9 and H1-4 being provided in SEQ ID NO: 1 and 4-7, respectively.

The cDNA sequences for the isolated clones were compared to known sequences in the gene bank using the EMBL and GenBank databases (release 96). Four of the prostate tumor cDNA clones, F1-13, F1-16, H1-1, and H1-4, were determined to encode the following previously identified proteins: prostate specific antigen (PSA), human glandular kallikrein, human tumor expression enhanced gene, and mitochondria cytochrome C oxidase subunit II. H1-9 was found to be identical to a previously identified human

autonomously replicating sequence. No significant homologies to the cDNA sequence for F1-12 were found.

Subsequent studies led to the isolation of a full-length cDNA sequence for F1-12. This sequence is provided in SEQ ID NO: 107, with the corresponding predicted amino acid sequence being provided in SEQ ID NO: 108.

To clone less abundant prostate tumor specific genes, cDNA library subtraction was performed by subtracting the prostate tumor cDNA library described above with the normal pancreas cDNA library and with the three most abundant genes in the previously subtracted prostate tumor specific cDNA library: human glandular kallikrein, prostate specific antigen (PSA), and mitochondria cytochrome C oxidase subunit II. Specifically, 1 µg each of human glandular kallikrein, PSA and mitochondria cytochrome C oxidase subunit II cDNAs in pCDNA3.1 were added to the driver DNA and subtraction was performed as described above to provide a second subtracted cDNA library hereinafter referred to as the "subtracted prostate tumor specific cDNA library with spike".

Twenty-two cDNA clones were isolated from the subtracted prostate tumor specific cDNA library with spike. The determined 3' and 5' cDNA sequences for the clones referred to as J1-17, L1-12, N1-1862, J1-13, J1-19, J1-25, J1-24, K1-58, K1-63, L1-4 and L1-14 are provided in SEQ ID NOS: 8-9, 10-11, 12-13, 14-15, 16-17, 18-19, 20-21, 22-23, 24-25, 26-27 and 28-29, respectively. The determined 3' cDNA sequences for the clones referred to as J1-12, J1-16, J1-21, K1-48, K1-55, L1-2, L1-6, N1-1858, N1-1860, N1-1861, N1-1864 are provided in SEQ ID NOS: 30-40, respectively. Comparison of these sequences with those in the gene bank as described above, revealed no significant homologies to three of the five most abundant DNA species, (J1-17, L1-12 and N1-1862; SEQ ID NOS: 8-9, 10-11 and 12-13, respectively). Of the remaining two most abundant species, one (J1-12; SEQ ID NO:30) was found to be identical to the previously identified human pulmonary surfactant-associated protein, and the other (K1-48; SEQ ID NO:33) was determined to have some homology to *R. norvegicus* mRNA for 2-arylpropionyl-CoA epimerase. Of the 17 less abundant cDNA clones isolated from the subtracted prostate tumor specific cDNA library with spike, four (J1-16, K1-55, L1-6 and N1-1864; SEQ ID NOS:31, 34, 36 and 40, respectively) were found to be identical to previously identified sequences, two (J1-21 and N1-1860; SEQ ID NOS: 32 and 38, respectively) were found to show some homology to non-human sequences, and two (L1-2 and N1-1861; SEQ ID NOS: 35 and 39, respectively) were found to show some homology to known human sequences. No significant homologies were found to the polypeptides J1-13, J1-19, J1-24, J1-25, K1-58, K1-63, L1-4, L1-14 (SEQ ID NOS: 14-15, 16-17, 20-21, 18-19, 22-23, 24-25, 26-27, 28-29, respectively).

Subsequent studies led to the isolation of full length cDNA sequences for J1-17, L1-12 and N1-1862 (SEQ ID NOS: 109-111, respectively). The corresponding predicted

amino acid sequences are provided in SEQ ID NOS: 112-114. L1-12 is also referred to as P501S.

In a further experiment, four additional clones were identified by subtracting a prostate tumor cDNA library with normal prostate cDNA prepared from a pool of three normal prostate poly A+ RNA (referred to as "prostate subtraction 2"). The determined cDNA sequences for these clones, hereinafter referred to as U1-3064, U1-3065, V1-3692 and 1A-3905, are provided in SEQ ID NO: 69-72, respectively. Comparison of the determined sequences with those in the gene bank revealed no significant homologies to U1-3065.

A second subtraction with spike (referred to as "prostate subtraction spike 2") was performed by subtracting a prostate tumor specific cDNA library with spike with normal pancreas cDNA library and further spiked with PSA, J1-17, pulmonary surfactant-associated protein, mitochondrial DNA, cytochrome c oxidase subunit II, N1-1862, autonomously replicating sequence, L1-12 and tumor expression enhanced gene. Four additional clones, hereinafter referred to as V1-3686, R1-2330, 1B-3976 and V1-3679, were isolated. The determined cDNA sequences for these clones are provided in SEQ ID NO: 73-76, respectively. Comparison of these sequences with those in the gene bank revealed no significant homologies to V1-3686 and R1-2330.

Further analysis of the three prostate subtractions described above (prostate subtraction 2, subtracted prostate tumor specific cDNA library with spike, and prostate subtraction spike 2) resulted in the identification of sixteen additional clones, referred to as 1G-4736, 1G-4738, 1G-4741, 1G-4744, 1G-4734, 1H-4774, 1H-4781, 1H-4785, 1H-4787, 1H-4796, 1I-4810, 1I-4811, 1J-4876, 1K-4884 and 1K-4896. The determined cDNA sequences for these clones are provided in SEQ ID NOS: 77-92, respectively. Comparison of these sequences with those in the gene bank as described above, revealed no significant homologies to 1G-4741, 1G-4734, 1I-4807, 1J-4876 and 1K-4896 (SEQ ID NOS: 79, 81, 87, 90 and 92, respectively). Further analysis of the isolated clones led to the determination of extended cDNA sequences for 1G-4736, 1G-4738, 1G-4741, 1G-4744, 1H-4774, 1H-4781, 1H-4785, 1H-4787, 1H-4796, 1I-4807, 1J-4876, 1K-4884 and 1K-4896, provided in SEQ ID NOS: 179-188 and 191-193, respectively, and to the determination of additional partial cDNA sequences for 1I-4810 and 1I-4811, provided in SEQ ID NOS: 189 and 190, respectively.

Additional studies with prostate subtraction spike 2 resulted in the isolation of three more clones. Their sequences were determined as described above and compared to the most recent GenBank. All three clones were found to have homology to known genes, which are Cysteine-rich protein, KIAA0242, and KIAA0280 (SEQ ID NO: 317, 319, and 320, respectively). Further analysis of these clones by Synteni microarray (Synteni, Palo Alto, CA) demonstrated that all three clones were over-expressed in most prostate tumors and

prostate BPH, as well as in the majority of normal prostate tissues tested, but low expression in all other normal tissues.

An additional subtraction was performed by subtracting a normal prostate cDNA library with normal pancreas cDNA (referred to as "prostate subtraction 3"). This led to the identification of six additional clones referred to as 1G-4761, 1G-4762, 1H-4766, 1H-4770, 1H-4771 and 1H-4772 (SEQ ID NOS: 93-98). Comparison of these sequences with those in the gene bank revealed no significant homologies to 1G-4761 and 1H-4771 (SEQ ID NOS: 93 and 97, respectively). Further analysis of the isolated clones led to the determination of extended cDNA sequences for 1G-4761, 1G-4762, 1H-4766 and 1H-4772 provided in SEQ ID NOS: 194-196 and 199, respectively, and to the determination of additional partial cDNA sequences for 1H-4770 and 1H-4771, provided in SEQ ID NOS: 197 and 198, respectively.

Subtraction of a prostate tumor cDNA library, prepared from a pool of polyA+ RNA from three prostate cancer patients, with a normal pancreas cDNA library (prostate subtraction 4) led to the identification of eight clones, referred to as 1D-4297, 1D-4309, 1D.1-4278, 1D-4288, 1D-4283, 1D-4304, 1D-4296 and 1D-4280 (SEQ ID NOS: 99-107). These sequences were compared to those in the gene bank as described above. No significant homologies were found to 1D-4283 and 1D-4304 (SEQ ID NOS: 103 and 104, respectively). Further analysis of the isolated clones led to the determination of extended cDNA sequences for 1D-4309, 1D.1-4278, 1D-4288, 1D-4283, 1D-4304, 1D-4296 and 1D-4280, provided in SEQ ID NOS: 200-206, respectively.

cDNA clones isolated in prostate subtraction 1 and prostate subtraction 2, described above, were colony PCR amplified and their mRNA expression levels in prostate tumor, normal prostate and in various other normal tissues were determined using microarray technology (Synteni, Palo Alto, CA). Briefly, the PCR amplification products were dotted onto slides in an array format, with each product occupying a unique location in the array. mRNA was extracted from the tissue sample to be tested, reverse transcribed, and fluorescent-labeled cDNA probes were generated. The microarrays were probed with the labeled cDNA probes, the slides scanned and fluorescence intensity was measured. This intensity correlates with the hybridization intensity. Two clones (referred to as P509S and P510S) were found to be over-expressed in prostate tumor and normal prostate and expressed at low levels in all other normal tissues tested (liver, pancreas, skin, bone marrow, brain, breast, adrenal gland, bladder, testes, salivary gland, large intestine, kidney, ovary, lung, spinal cord, skeletal muscle and colon). The determined cDNA sequences for P509S and P510S are provided in SEQ ID NO: 223 and 224, respectively. Comparison of these sequences with those in the gene bank as described above, revealed some homology to previously identified ESTs.

Additional, studies led to the isolation of the full-length cDNA sequence for P509S. This sequence is provided in SEQ ID NO: 332, with the corresponding predicted amino acid sequence being provided in SEQ ID NO: 339.

## EXAMPLE 2

### DETERMINATION OF TISSUE SPECIFICITY OF PROSTATE TUMOR POLYPEPTIDES

Using gene specific primers, mRNA expression levels for the representative prostate tumor polypeptides F1-16, H1-1, J1-17 (also referred to as P502S), L1-12 (also referred to as P501S), F1-12 (also referred to as P504S) and N1-1862 (also referred to as P503S) were examined in a variety of normal and tumor tissues using RT-PCR.

Briefly, total RNA was extracted from a variety of normal and tumor tissues using Trizol reagent as described above. First strand synthesis was carried out using 1-2  $\mu$ g of total RNA with SuperScript II reverse transcriptase (BRL Life Technologies) at 42 °C for one hour. The cDNA was then amplified by PCR with gene-specific primers. To ensure the semi-quantitative nature of the RT-PCR,  $\beta$ -actin was used as an internal control for each of the tissues examined. First, serial dilutions of the first strand cDNAs were prepared and RT-PCR assays were performed using  $\beta$ -actin specific primers. A dilution was then chosen that enabled the linear range amplification of the  $\beta$ -actin template and which was sensitive enough to reflect the differences in the initial copy numbers. Using these conditions, the  $\beta$ -actin levels were determined for each reverse transcription reaction from each tissue. DNA contamination was minimized by DNase treatment and by assuring a negative PCR result when using first strand cDNA that was prepared without adding reverse transcriptase.

mRNA Expression levels were examined in four different types of tumor tissue (prostate tumor from 2 patients, breast tumor from 3 patients, colon tumor, lung tumor), and sixteen different normal tissues, including prostate, colon, kidney, liver, lung, ovary, pancreas, skeletal muscle, skin, stomach, testes, bone marrow and brain. F1-16 was found to be expressed at high levels in prostate tumor tissue, colon tumor and normal prostate, and at lower levels in normal liver, skin and testes, with expression being undetectable in the other tissues examined. H1-1 was found to be expressed at high levels in prostate tumor, lung tumor, breast tumor, normal prostate, normal colon and normal brain, at much lower levels in normal lung, pancreas, skeletal muscle, skin, small intestine, bone marrow, and was not detected in the other tissues tested. J1-17 (P502S) and L1-12 (P501S) appear to be specifically over-expressed in prostate, with both genes being expressed at high levels in prostate tumor and normal prostate but at low to undetectable levels in all the other tissues examined. N1-1862 (P503S) was found to be over-expressed in 60% of prostate tumors and detectable in normal colon and kidney. The RT-PCR results thus indicate that

F1-16, H1-1, J1-17 (P502S), N1-1862 (P503S) and L1-12 (P501S) are either prostate specific or are expressed at significantly elevated levels in prostate.

Further RT-PCR studies showed that F1-12 (P504S) is over-expressed in 60% of prostate tumors, detectable in normal kidney but not detectable in all other tissues tested. Similarly, R1-2330 was shown to be over-expressed in 40% of prostate tumors, detectable in normal kidney and liver, but not detectable in all other tissues tested. U1-3064 was found to be over-expressed in 60% of prostate tumors, and also expressed in breast and colon tumors, but was not detectable in normal tissues.

RT-PCR characterization of R1-2330, U1-3064 and 1D-4279 showed that these three antigens are over-expressed in prostate and/or prostate tumors.

Northern analysis with four prostate tumors, two normal prostate samples, two BPH prostates, and normal colon, kidney, liver, lung, pancreas, skeletal muscle, brain, stomach, testes, small intestine and bone marrow, showed that L1-12 (P501S) is over-expressed in prostate tumors and normal prostate, while being undetectable in other normal tissues tested. J1-17 (P502S) was detected in two prostate tumors and not in the other tissues tested. N1-1862 (P503S) was found to be over-expressed in three prostate tumors and to be expressed in normal prostate, colon and kidney, but not in other tissues tested. F1-12 (P504S) was found to be highly expressed in two prostate tumors and to be undetectable in all other tissues tested.

The microarray technology described above was used to determine the expression levels of representative antigens described herein in prostate tumor, breast tumor and the following normal tissues: prostate, liver, pancreas, skin, bone marrow, brain, breast, adrenal gland, bladder, testes, salivary gland, large intestine, kidney, ovary, lung, spinal cord, skeletal muscle and colon. L1-12 (P501S) was found to be over-expressed in normal prostate and prostate tumor, with some expression being detected in normal skeletal muscle. Both J1-12 and F1-12 (P504S) were found to be over-expressed in prostate tumor, with expression being lower or undetectable in all other tissues tested. N1-1862 (P503S) was found to be expressed at high levels in prostate tumor and normal prostate, and at low levels in normal large intestine and normal colon, with expression being undetectable in all other tissues tested. R1-2330 was found to be over-expressed in prostate tumor and normal prostate, and to be expressed at lower levels in all other tissues tested. 1D-4279 was found to be over-expressed in prostate tumor and normal prostate, expressed at lower levels in normal spinal cord, and to be undetectable in all other tissues tested.

Further microarray analysis to specifically address the extent to which P501S (SEQ ID NO: 110) was expressed in breast tumor revealed moderate over-expression not only in breast tumor, but also in metastatic breast tumor (2/31), with negligible to low expression

in normal tissues. This data suggests that P501S may be over-expressed in various breast tumors as well as in prostate tumors.

The expression levels of 32 ESTs (expressed sequence tags) described by Vasmatzis *et al.* (*Proc. Natl. Acad. Sci. USA* 95:300-304, 1998) in a variety of tumor and normal tissues were examined by microarray technology as described above. Two of these clones (referred to as P1000C and P1001C) were found to be over-expressed in prostate tumor and normal prostate, and expressed at low to undetectable levels in all other tissues tested (normal aorta, thymus, resting and activated PBMC, epithelial cells, spinal cord, adrenal gland, fetal tissues, skin, salivary gland, large intestine, bone marrow, liver, lung, dendritic cells, stomach, lymph nodes, brain, heart, small intestine, skeletal muscle, colon and kidney. The determined cDNA sequences for P1000C and P1001C are provided in SEQ ID NO: 384 and 472, respectively. The sequence of P1001C was found to show some homology to the previously isolated Human mRNA for JM27 protein. No significant homologies were found to the sequence of P1000C.

The expression of the polypeptide encoded by the full length cDNA sequence for F1-12 (also referred to as P504S; SEQ ID NO: 108) was investigated by immunohistochemical analysis. Rabbit-anti-P504S polyclonal antibodies were generated against the full length P504S protein by standard techniques. Subsequent isolation and characterization of the polyclonal antibodies were also performed by techniques well known in the art. Immunohistochemical analysis showed that the P504S polypeptide was expressed in 100% of prostate carcinoma samples tested (n=5).

The rabbit-anti-P504S polyclonal antibody did not appear to label benign prostate cells with the same cytoplasmic granular staining, but rather with light nuclear staining. Analysis of normal tissues revealed that the encoded polypeptide was found to be expressed in some, but not all normal human tissues. Positive cytoplasmic staining with rabbit-anti-P504S polyclonal antibody was found in normal human kidney, liver, brain, colon and lung-associated macrophages, whereas heart and bone marrow were negative.

This data indicates that the P504S polypeptide is present in prostate cancer tissues, and that there are qualitative and quantitative differences in the staining between benign prostatic hyperplasia tissues and prostate cancer tissues, suggesting that this polypeptide may be detected selectively in prostate tumors and therefore be useful in the diagnosis of prostate cancer.

### EXAMPLE 3

#### ISOLATION AND CHARACTERIZATION OF PROSTATE TUMOR POLYPEPTIDES BY PCR-BASED SUBTRACTION

A cDNA subtraction library, containing cDNA from normal prostate subtracted with ten other normal tissue cDNAs (brain, heart, kidney, liver, lung, ovary, placenta, skeletal muscle, spleen and thymus) and then submitted to a first round of PCR amplification, was purchased from Clontech. This library was subjected to a second round of PCR amplification, following the manufacturer's protocol. The resulting cDNA fragments were subcloned into the vector pT7 Blue T-vector (Novagen, Madison, WI) and transformed into XL-1 Blue MRF' *E. coli* (Stratagene). DNA was isolated from independent clones and sequenced using a Perkin Elmer/Applied Biosystems Division Automated Sequencer Model 373A.

Fifty-nine positive clones were sequenced. Comparison of the DNA sequences of these clones with those in the gene bank, as described above, revealed no significant homologies to 25 of these clones, hereinafter referred to as P5, P8, P9, P18, P20, P30, P34, P36, P38, P39, P42, P49, P50, P53, P55, P60, P64, P65, P73, P75, P76, P79 and P84. The determined cDNA sequences for these clones are provided in SEQ ID NO: 41-45, 47-52 and 54-65, respectively. P29, P47, P68, P80 and P82 (SEQ ID NO: 46, 53 and 66-68, respectively) were found to show some degree of homology to previously identified DNA sequences. To the best of the inventors' knowledge, none of these sequences have been previously shown to be present in prostate.

Further studies using the PCR-based methodology described above resulted in the isolation of more than 180 additional clones, of which 23 clones were found to show no significant homologies to known sequences. The determined cDNA sequences for these clones are provided in SEQ ID NO: 115-123, 127, 131, 137, 145, 147-151, 153, 156-158 and 160. Twenty-three clones (SEQ ID NO: 124-126, 128-130, 132-136, 138-144, 146, 152, 154, 155 and 159) were found to show some homology to previously identified ESTs. An additional ten clones (SEQ ID NO: 161-170) were found to have some degree of homology to known genes. Larger cDNA clones containing the P20 sequence represent splice variants of a gene referred to as P703P. The determined DNA sequence for the variants referred to as DE1, DE13 and DE14 are provided in SEQ ID NOS: 171, 175 and 177, respectively, with the corresponding predicted amino acid sequences being provided in SEQ ID NO: 172, 176 and 178, respectively. The determined cDNA sequence for an extended spliced form of P703 is provided in SEQ ID NO: 225. The DNA sequences for the splice variants referred to as DE2 and DE6 are provided in SEQ ID NOS: 173 and 174, respectively.

mRNA Expression levels for representative clones in tumor tissues (prostate (n=5), breast (n=2), colon and lung) normal tissues (prostate (n=5), colon, kidney, liver, lung (n=2), ovary (n=2), skeletal muscle, skin, stomach, small intestine and brain), and activated



and non-activated PBMC was determined by RT-PCR as described above. Expression was examined in one sample of each tissue type unless otherwise indicated.

P9 was found to be highly expressed in normal prostate and prostate tumor compared to all normal tissues tested except for normal colon which showed comparable expression. P20, a portion of the P703P gene, was found to be highly expressed in normal prostate and prostate tumor, compared to all twelve normal tissues tested. A modest increase in expression of P20 in breast tumor (n=2), colon tumor and lung tumor was seen compared to all normal tissues except lung (1 of 2). Increased expression of P18 was found in normal prostate, prostate tumor and breast tumor compared to other normal tissues except lung and stomach. A modest increase in expression of P5 was observed in normal prostate compared to most other normal tissues. However, some elevated expression was seen in normal lung and PBMC. Elevated expression of P5 was also observed in prostate tumors (2 of 5), breast tumor and one lung tumor sample. For P30, similar expression levels were seen in normal prostate and prostate tumor, compared to six of twelve other normal tissues tested. Increased expression was seen in breast tumors, one lung tumor sample and one colon tumor sample, and also in normal PBMC. P29 was found to be over-expressed in prostate tumor (5 of 5) and normal prostate (5 of 5) compared to the majority of normal tissues. However, substantial expression of P29 was observed in normal colon and normal lung (2 of 2). P80 was found to be over-expressed in prostate tumor (5 of 5) and normal prostate (5 of 5) compared to all other normal tissues tested, with increased expression also being seen in colon tumor.

Further studies resulted in the isolation of twelve additional clones, hereinafter referred to as 10-d8, 10-h10, 11-c8, 7-g6, 8-b5, 8-b6, 8-d4, 8-d9, 8-g3, 8-h11, 9-f12 and 9-f3. The determined DNA sequences for 10-d8, 10-h10, 11-c8, 8-d4, 8-d9, 8-h11, 9-f12 and 9-f3 are provided in SEQ ID NO: 207, 208, 209, 216, 217, 220, 221 and 222, respectively. The determined forward and reverse DNA sequences for 7-g6, 8-b5, 8-b6 and 8-g3 are provided in SEQ ID NO: 210 and 211; 212 and 213; 214 and 215; and 218 and 219, respectively. Comparison of these sequences with those in the gene bank revealed no significant homologies to the sequence of 9-f3. The clones 10-d8, 11-c8 and 8-h11 were found to show some homology to previously isolated ESTs, while 10-h10, 8-b5, 8-b6, 8-d4, 8-d9, 8-g3 and 9-f12 were found to show some homology to previously identified genes. Further characterization of 7-G6 and 8-G3 showed identity to the known genes PAP and PSA, respectively.

mRNA expression levels for these clones were determined using the microarray technology described above. The clones 7-G6, 8-G3, 8-B5, 8-B6, 8-D4, 8-D9, 9-F3, 9-F12, 9-H3, 10-A2, 10-A4, 11-C9 and 11-F2 were found to be over-expressed in prostate tumor and normal prostate, with expression in other tissues tested being low or undetectable.

Increased expression of 8-F11 was seen in prostate tumor and normal prostate, bladder, skeletal muscle and colon. Increased expression of 10-H10 was seen in prostate tumor and normal prostate, bladder, lung, colon, brain and large intestine. Increased expression of 9-B1 was seen in prostate tumor, breast tumor, and normal prostate, salivary gland, large intestine and skin, with increased expression of 11-C8 being seen in prostate tumor, and normal prostate and large intestine.

An additional cDNA fragment derived from the PCR-based normal prostate subtraction, described above, was found to be prostate specific by both micro-array technology and RT-PCR. The determined cDNA sequence of this clone (referred to as 9-A11) is provided in SEQ ID NO: 226. Comparison of this sequence with those in the public databases revealed 99% identity to the known gene HOXB13.

Further studies led to the isolation of the clones 8-C6 and 8-H7. The determined cDNA sequences for these clones are provided in SEQ ID NO: 227 and 228, respectively. These sequences were found to show some homology to previously isolated ESTs.

PCR and hybridization-based methodologies were employed to obtain longer cDNA sequences for clone P20 (also referred to as P703P), yielding three additional cDNA fragments that progressively extend the 5' end of the gene. These fragments, referred to as P703PDE5, P703P6.26, and P703PX-23 (SEQ ID NO: 326, 328 and 330, with the predicted corresponding amino acid sequences being provided in SEQ ID NO: 327, 329 and 331, respectively) contain additional 5' sequence. P703PDE5 was recovered by screening of a cDNA library (#141-26) with a portion of P703P as a probe. P703P6.26 was recovered from a mixture of three prostate tumor cDNAs and P703PX\_23 was recovered from cDNA library (#438-48). Together, the additional sequences include all of the putative mature serine protease along with part of the putative signal sequence. Further studies using a PCR-based subtraction library of a prostate tumor pool subtracted against a pool of normal tissues (referred to as JP: PCR subtraction) resulted in the isolation of thirteen additional clones, seven of which did not share any significant homology to known GenBank sequences. The determined cDNA sequences for these seven clones (P711P, P712P, novel 23, P774P, P775P, P710P and P768P) are provided in SEQ ID NO: 307-311, 313 and 315, respectively. The remaining six clones (SEQ ID NO: 316 and 321-325) were shown to share some homology to known genes. By microarray analysis, all thirteen clones showed three or more fold over-expression in prostate tissues, including prostate tumors, BPH and normal prostate as compared to normal non-prostate tissues. Clones P711P, P712P, novel 23 and P768P showed over-expression in most prostate tumors and BPH tissues tested (n=29), and in the majority of normal prostate tissues (n=4), but background to low expression levels in all normal tissues.

Clones P774P, P775P and P710P showed comparatively lower expression and expression in fewer prostate tumors and BPH samples, with negative to low expression in normal prostate.

The full-length cDNA for P711P was obtained by employing the partial sequence of SEQ ID NO: 307 to screen a prostate cDNA library. Specifically, a directionally cloned prostate cDNA library was prepared using standard techniques. One million colonies of this library were plated onto LB/Amp plates. Nylon membrane filters were used to lift these colonies, and the cDNAs which were picked up by these filters were denatured and cross-linked to the filters by UV light. The P711P cDNA fragment of SEQ ID NO: 307 was radio-labeled and used to hybridize with these filters. Positive clones were selected, and cDNAs were prepared and sequenced using an automatic Perkin Elmer/Applied Biosystems sequencer. The determined full-length sequence of P711P is provided in SEQ ID NO: 382, with the corresponding predicted amino acid sequence being provided in SEQ ID NO: 383.

Using PCR and hybridization-based methodologies, additional cDNA sequence information was derived for two clones described above, 11-C9 and 9-F3, herein after referred to as P707P and P714P, respectively (SEQ ID NO: 333 and 334). After comparison with the most recent GenBank, P707P was found to be a splice variant of the known gene HoxB13. In contrast, no significant homologies to P714P were found.

Clones 8-B3, P89, P98, P130 and P201 (as disclosed in U.S. Patent Application No. 09/020,956, filed February 9, 1998) were found to be contained within one contiguous sequence, referred to as P705P (SEQ ID NO: 335, with the predicted amino acid sequence provided in SEQ ID NO: 336), which was determined to be a splice variant of the known gene NKX 3.1.

#### EXAMPLE 4 SYNTHESIS OF POLYPEPTIDES

Polypeptides may be synthesized on a Perkin Elmer/Applied Biosystems 430A peptide synthesizer using Fmoc chemistry with HPTU (O-Benzotriazole-N,N,N',N'-tetramethyluronium hexafluorophosphate) activation. A Gly-Cys-Gly sequence may be attached to the amino terminus of the peptide to provide a method of conjugation, binding to an immobilized surface, or labeling of the peptide. Cleavage of the peptides from the solid support may be carried out using the following cleavage mixture: trifluoroacetic acid:ethanedithiol:thioanisole:water:phenol (40:1:2:2:3). After cleaving for 2 hours, the peptides may be precipitated in cold methyl-t-butyl-ether. The peptide pellets may then be dissolved in water containing 0.1% trifluoroacetic acid (TFA) and lyophilized prior to purification by C18 reverse phase HPLC. A gradient of 0%-60% acetonitrile (containing 0.1% TFA) in water (containing 0.1% TFA) may be used to elute the peptides. Following

lyophilization of the pure fractions, the peptides may be characterized using electrospray or other types of mass spectrometry and by amino acid analysis.

#### EXAMPLE 5

#### FURTHER ISOLATION AND CHARACTERIZATION OF PROSTATE TUMOR POLYPEPTIDES BY PCR-BASED SUBTRACTION

A cDNA library generated from prostate primary tumor mRNA as described above was subtracted with cDNA from normal prostate. The subtraction was performed using a PCR-based protocol (Clontech), which was modified to generate larger fragments. Within this protocol, tester and driver double stranded cDNA were separately digested with five restriction enzymes that recognize six-nucleotide restriction sites (MluI, MscI, PvuII, SalI and StuI). This digestion resulted in an average cDNA size of 600 bp, rather than the average size of 300 bp that results from digestion with RsaI according to the Clontech protocol. This modification did not affect the subtraction efficiency. Two tester populations were then created with different adapters, and the driver library remained without adapters.

The tester and driver libraries were then hybridized using excess driver cDNA. In the first hybridization step, driver was separately hybridized with each of the two tester cDNA populations. This resulted in populations of (a) unhybridized tester cDNAs, (b) tester cDNAs hybridized to other tester cDNAs, (c) tester cDNAs hybridized to driver cDNAs and (d) unhybridized driver cDNAs. The two separate hybridization reactions were then combined, and rehybridized in the presence of additional denatured driver cDNA. Following this second hybridization, in addition to populations (a) through (d), a fifth population (e) was generated in which tester cDNA with one adapter hybridized to tester cDNA with the second adapter. Accordingly, the second hybridization step resulted in enrichment of differentially expressed sequences which could be used as templates for PCR amplification with adaptor-specific primers.

The ends were then filled in, and PCR amplification was performed using adaptor-specific primers. Only population (e), which contained tester cDNA that did not hybridize to driver cDNA, was amplified exponentially. A second PCR amplification step was then performed, to reduce background and further enrich differentially expressed sequences.

This PCR-based subtraction technique normalizes differentially expressed cDNAs so that rare transcripts that are overexpressed in prostate tumor tissue may be recoverable. Such transcripts would be difficult to recover by traditional subtraction methods.

In addition to genes known to be overexpressed in prostate tumor, seventy-seven further clones were identified. Sequences of these partial cDNAs are provided in SEQ ID NO: 29 to 305. Most of these clones had no significant homology to database sequences. Exceptions were JTPN23 (SEQ ID NO: 231; similarity to pig valosin-containing protein), JTPN30 (SEQ ID NO: 234; similarity to rat mRNA for proteasome subunit), JTPN45 (SEQ ID NO: 243; similarity to rat *norvegicus* cytosolic NADP-dependent isocitrate dehydrogenase), JTPN46 (SEQ ID NO: 244; similarity to human subclone H8 4 d4 DNA sequence), JP1D6 (SEQ ID NO: 265; similarity to *G. gallus* dynein light chain-A), JP8D6 (SEQ ID NO: 288; similarity to human BAC clone RG016J04), JP8F5 (SEQ ID NO: 289; similarity to human subclone H8 3 b5 DNA sequence), and JP8E9 (SEQ ID NO: 299; similarity to human Alu sequence).

Additional studies using the PCR-based subtraction library consisting of a prostate tumor pool subtracted against a normal prostate pool (referred to as PT-PN PCR subtraction) yielded three additional clones. Comparison of the cDNA sequences of these clones with the most recent release of GenBank revealed no significant homologies to the two clones referred to as P715P and P767P (SEQ ID NO: 312 and 314). The remaining clone was found to show some homology to the known gene KIAA0056 (SEQ ID NO: 318). Using microarray analysis to measure mRNA expression levels in various tissues, all three clones were found to be over-expressed in prostate tumors and BPH tissues. Specifically, clone P715P was over-expressed in most prostate tumors and BPH tissues by a factor of three or greater, with elevated expression seen in the majority of normal prostate samples and in fetal tissue, but negative to low expression in all other normal tissues. Clone P767P was over-expressed in several prostate tumors and BPH tissues, with moderate expression levels in half of the normal prostate samples, and background to low expression in all other normal tissues tested.

Further analysis, by microarray as described above, of the PT-PN PCR subtraction library and of a DNA subtraction library containing cDNA from prostate tumor subtracted with a pool of normal tissue cDNAs, led to the isolation of 27 additional clones (SEQ ID NO: 340-365 and 381) which were determined to be over-expressed in prostate tumor. The clones of SEQ ID NO: 341, 342, 345, 347, 348, 349, 351, 355-359, 361, 362 and 364 were also found to be expressed in normal prostate. Expression of all 26 clones in a variety of normal tissues was found to be low or undetectable, with the exception of P544S (SEQ ID NO: 356) which was found to be expressed in small intestine. Of the 26 clones, 10 (SEQ ID NO: 340-349) were found to show some homology to previously identified sequences. No significant homologies were found to the clones of SEQ ID NO: 350-365.

#### EXAMPLE 6

##### PEPTIDE PRIMING OF MICE AND PROPAGATION OF CTL LINES

6.1. This Example illustrates the preparation of a CTL cell line specific for cells expressing the P502S gene.

Mice expressing the transgene for human HLA A2.1 (provided by Dr L. Sherman, The Scripps Research Institute, La Jolla, CA) were immunized with P2S#12 peptide (VLGWVAEL; SEQ ID NO: 306), which is derived from the P502S gene (also referred to herein as J1-17, SEQ ID NO: 8), as described by Theobald et al., *Proc. Natl. Acad. Sci. USA* 92:11993-11997, 1995 with the following modifications. Mice were immunized with 100µg of P2S#12 and 120µg of an I-A<sup>b</sup> binding peptide derived from hepatitis B Virus protein emulsified in incomplete Freund's adjuvant. Three weeks later these mice were sacrificed and using a nylon mesh single cell suspensions prepared. Cells were then resuspended at  $6 \times 10^6$  cells/ml in complete media (RPMI-1640; Gibco BRL, Gaithersburg, MD) containing 10% FCS, 2mM Glutamine (Gibco BRL), sodium pyruvate (Gibco BRL), non-essential amino acids (Gibco BRL),  $2 \times 10^{-5}$  M 2-mercaptoethanol, 50U/ml penicillin and streptomycin, and cultured in the presence of irradiated (3000 rads) P2S#12-pulsed (5mg/ml P2S#12 and 10mg/ml  $\beta$ 2-microglobulin) LPS blasts (A2 transgenic spleens cells cultured in the presence of 7µg/ml dextran sulfate and 25µg/ml LPS for 3 days). Six days later, cells ( $5 \times 10^5$ /ml) were restimulated with  $2.5 \times 10^6$ /ml peptide pulsed irradiated (20,000 rads) EL4A2Kb cells (Sherman et al, *Science* 258:815-818, 1992) and  $3 \times 10^6$ /ml A2 transgenic spleen feeder cells. Cells were cultured in the presence of 20U/ml IL-2. Cells continued to be restimulated on a weekly basis as described, in preparation for cloning the line.

P2S#12 line was cloned by limiting dilution analysis with peptide pulsed EL4 A2Kb tumor cells ( $1 \times 10^4$  cells/ well) as stimulators and A2 transgenic spleen cells as feeders ( $5 \times 10^5$  cells/ well) grown in the presence of 30U/ml IL-2. On day 14, cells were

restimulated as before. On day 21, clones that were growing were isolated and maintained in culture. Several of these clones demonstrated significantly higher reactivity (lysis) against human fibroblasts (HLA A2.1 expressing) transduced with P502S than against control fibroblasts. An example is presented in Figure 1.

This data indicates that P2S #12 represents a naturally processed epitope of the P502S protein that is expressed in the context of the human HLA A2.1 molecule.

6.2. This Example illustrates the preparation of murine CTL lines and CTL clones specific for cells expressing the P501S gene.

This series of experiments were performed similarly to that described above. Mice were immunized with the P1S#10 peptide (SEQ ID NO: 337), which is derived from the P501S gene (also referred to herein as L1-12, SEQ ID NO: 110). The P1S#10 peptide was derived by analysis of the predicted polypeptide sequence for P501S for potential HLA-A2 binding sequences as defined by published HLA-A2 binding motifs (Parker, KC, *et al*, *J. Immunol.*, 152:163, 1994). P1S#10 peptide was synthesized as described in Example 4, and empirically tested for HLA-A2 binding using a T cell based competition assay. Predicted A2 binding peptides were tested for their ability to compete HLA-A2 specific peptide presentation to an HLA-A2 restricted CTL clone (D150M58), which is specific for the HLA-A2 binding influenza matrix peptide fluM58. D150M58 CTL secretes TNF in response to self-presentation of peptide fluM58. In the competition assay, test peptides at 100-200  $\mu\text{g/ml}$  were added to cultures of D150M58 CTL in order to bind HLA-A2 on the CTL. After thirty minutes, CTL cultured with test peptides, or control peptides, were tested for their antigen dose response to the fluM58 peptide in a standard TNF bioassay. As shown in Figure 3, peptide P1S#10 competes HLA-A2 restricted presentation of fluM58, demonstrating that peptide P1S#10 binds HLA-A2.

Mice expressing the transgene for human HLA A2.1 were immunized as described by Theobald *et al.* (*Proc. Natl. Acad. Sci. USA* 92:11993-11997, 1995) with the following modifications. Mice were immunized with 62.5 $\mu\text{g}$  of P1S #10 and 120 $\mu\text{g}$  of an I-A<sup>b</sup> binding peptide derived from Hepatitis B Virus protein emulsified in incomplete Freund's adjuvant. Three weeks later these mice were sacrificed and single cell suspensions prepared using a nylon mesh. Cells were then resuspended at  $6 \times 10^6$  cells/ml in complete media (as described above) and cultured in the presence of irradiated (3000 rads) P1S#10-pulsed (2 $\mu\text{g/ml}$  P1S#10 and 10mg/ml  $\beta$ 2-microglobulin) LPS blasts (A2 transgenic spleens cells cultured in the presence of 7 $\mu\text{g/ml}$  dextran sulfate and 25 $\mu\text{g/ml}$  LPS for 3 days). Six days later cells ( $5 \times 10^5$ /ml) were restimulated with  $2.5 \times 10^6$ /ml peptide-pulsed irradiated (20,000 rads) EL4A2Kb cells, as described above, and  $3 \times 10^6$ /ml A2 transgenic spleen feeder cells. Cells were cultured in the presence of 20 U/ml IL-2. Cells were restimulated on a weekly

basis in preparation for cloning. After three rounds of *in vitro* stimulations, one line was generated that recognized P1S#10-pulsed Jurkat A2Kb targets and P501S-transduced Jurkat targets as shown in Figure 4.

A P1S#10-specific CTL line was cloned by limiting dilution analysis with peptide pulsed EL4 A2Kb tumor cells ( $1 \times 10^4$  cells/ well) as stimulators and A2 transgenic spleen cells as feeders ( $5 \times 10^5$  cells/ well) grown in the presence of 30U/ml IL-2. On day 14, cells were restimulated as before. On day 21, viable clones were isolated and maintained in culture. As shown in Figure 5, five of these clones demonstrated specific cytolytic reactivity against P501S-transduced Jurkat A2Kb targets. This data indicates that P1S#10 represents a naturally processed epitope of the P501S protein that is expressed in the context of the human HLA-A2.1 molecule.

#### EXAMPLE 7

#### ABILITY OF HUMAN T CELLS TO RECOGNIZE PROSTATE TUMOR POLYPEPTIDES

This Example illustrates the ability of T cells specific for a prostate tumor polypeptide to recognize human tumor.

Human CD8<sup>+</sup> T cells were primed *in vitro* to the P2S-12 peptide (SEQ ID NO: 306) derived from P502S (also referred to as J1-17) using dendritic cells according to the protocol of Van Tsai et al. (*Critical Reviews in Immunology* 18:65-75, 1998). The resulting CD8<sup>+</sup> T cell microcultures were tested for their ability to recognize the P2S-12 peptide presented by autologous fibroblasts or fibroblasts which were transduced to express the P502S gene in a  $\gamma$ -interferon ELISPOT assay (see Lalvani et al., *J. Exp. Med.* 186:859-865, 1997). Briefly, titrating numbers of T cells were assayed in duplicate on  $10^4$  fibroblasts in the presence of 3  $\mu$ g/ml human  $\beta_2$ -microglobulin and 1  $\mu$ g/ml P2S-12 peptide or control E75 peptide. In addition, T cells were simultaneously assayed on autologous fibroblasts transduced with the P502S gene or as a control, fibroblasts transduced with HER-2/*neu*. Prior to the assay, the fibroblasts were treated with 10 ng/ml  $\gamma$ -interferon for 48 hours to upregulate class I MHC expression. One of the microcultures (#5) demonstrated strong recognition of both peptide pulsed fibroblasts as well as transduced fibroblasts in a  $\gamma$ -interferon ELISPOT assay. Figure 2A demonstrates that there was a strong increase in the number of  $\gamma$ -interferon spots with increasing numbers of T cells on fibroblasts pulsed with the P2S-12 peptide (solid bars) but not with the control E75 peptide (open bars). This shows the ability of these T cells to specifically recognize the P2S-12 peptide. As shown in Figure 2B, this microculture also demonstrated an increase in the number of  $\gamma$ -interferon spots with increasing numbers of T



cells on fibroblasts transduced to express the P502S gene but not the HER-2/*neu* gene. These results provide additional confirmatory evidence that the P2S-12 peptide is a naturally processed epitope of the P502S protein. Furthermore, this also demonstrates that there exists in the human T cell repertoire, high affinity T cells which are capable of recognizing this epitope. These T cells should also be capable of recognizing human tumors which express the P502S gene.

#### EXAMPLE 8

##### PRIMING OF CTL *IN VIVO* USING NAKED DNA IMMUNIZATION WITH A PROSTATE ANTIGEN

The prostate tumor antigen L1-12, as described above, is also referred to as P501S. HLA A2Kb Tg mice (provided by Dr L. Sherman, The Scripps Research Institute, La Jolla, CA) were immunized with 100 µg VR10132-P501S either intramuscularly or intradermally. The mice were immunized three times, with a two week interval between immunizations. Two weeks after the last immunization, immune spleen cells were cultured with Jurkat A2Kb-P501S transduced stimulator cells. CTL lines were stimulated weekly. After two weeks of *in vitro* stimulation, CTL activity was assessed against P501S transduced targets. Two out of 8 mice developed strong anti-P501S CTL responses. These results demonstrate that P501S contains at least one naturally processed A2-restricted CTL epitope.

#### EXAMPLE 9

##### GENERATION OF HUMAN CTL *IN VITRO* USING WHOLE GENE PRIMING AND STIMULATION TECHNIQUES WITH PROSTATE TUMOR ANTIGEN

Using *in vitro* whole-gene priming with P501S-retrovirally transduced autologous fibroblasts (see, for example, Yee et al, *The Journal of Immunology*, 157(9):4079-86, 1996), human CTL lines were derived that specifically recognize autologous fibroblasts transduced with P501S (also known as L1-12), as determined by interferon-γ ELISPOT analysis as described above. Using a panel of HLA-mismatched fibroblast lines transduced with P501S, these CTL lines were shown to be restricted HLA-A2 class I allele. Specifically, dendritic cells (DC) were differentiated from monocyte cultures derived from PBMC of normal human donors by growing for five days in RPMI medium containing 10% human serum, 50 ng/ml human GM-CSF and 30 ng/ml human IL-4. Following culture, DC were infected overnight with recombinant P501S vaccinia virus at a multiplicity of infection (M.O.I) of five, and matured overnight by the addition of 3 µg/ml CD40 ligand. Virus was inactivated by UV irradiation. CD8+ T cells were isolated using a magnetic bead system, and

priming cultures were initiated using standard culture techniques. Cultures were restimulated every 7-10 days using autologous primary fibroblasts retrovirally transduced with P501S. Following four stimulation cycles, CD8+ T cell lines were identified that specifically produced interferon- $\gamma$  when stimulated with P501S-transduced autologous fibroblasts. The P501S-specific activity could be sustained by the continued stimulation of the cultures with P501S-transduced fibroblasts in the presence of IL-15. A panel of HLA-mismatched fibroblast lines transduced with P501S were generated to define the restriction allele of the response. By measuring interferon- $\gamma$  in an ELISPOT assay, the P501S specific response was shown to be restricted by HLA-A2. These results demonstrate that a CD8+ CTL response to P501S can be elicited.

#### EXAMPLE 10

##### IDENTIFICATION OF A NATURALLY PROCESSED CTL EPITOPE CONTAINED WITHIN A PROSTATE TUMOR ANTIGEN

The 9-mer peptide p5 (SEQ ID NO: 338) was derived from the P703P antigen (also referred to as P20). The p5 peptide is immunogenic in human HLA-A2 donors and is a naturally processed epitope. Antigen specific CD8+ T cells can be primed following repeated *in vitro* stimulations with monocytes pulsed with p5 peptide. These CTL specifically recognize p5-pulsed target cells in both ELISPOT (as described above) and chromium release assays. Additionally, immunization of HLA-A2 transgenic mice with p5 leads to the generation of CTL lines which recognize a variety of P703P transduced target cells expressing either HLA-A2Kb or HLA-A2. Specifically, HLA-A2 transgenic mice were immunized subcutaneously in the footpad with 100  $\mu$ g of p5 peptide together with 140  $\mu$ g of hepatitis B virus core peptide (a Th peptide) in Freund's incomplete adjuvant. Three weeks post immunization, spleen cells from immunized mice were stimulated *in vitro* with peptide-pulsed LPS blasts. CTL activity was assessed by chromium release assay five days after primary *in vitro* stimulation. Retrovirally transduced cells expressing the control antigen P703P and HLA-A2Kb were used as targets. CTL lines that specifically recognized both p5-pulsed targets as well as P703P-expressing targets were identified.

Human *in vitro* priming experiments demonstrated that the p5 peptide is immunogenic in humans. Dendritic cells (DC) were differentiated from monocyte cultures derived from PBMC of normal human donors by culturing for five days in RPMI medium containing 10% human serum, 50 ng/ml human GM-CSF and 30 ng/ml human IL-4. Following culture, the DC were pulsed with p5 peptide and cultured with GM-CSF and IL-4 together with CD8+ T cell enriched PBMC. CTL lines were restimulated on a weekly basis

with p5-pulsed monocytes. Five to six weeks after initiation of the CTL cultures, CTL recognition of p5-pulsed target cells was demonstrated.

#### EXAMPLE 11 EXPRESSION OF A BREAST TUMOR-DERIVED ANTIGEN IN PROSTATE

Isolation of the antigen B305D from breast tumor by differential display is described in US Patent Application No. 08/700,014, filed August 20, 1996. Several different splice forms of this antigen were isolated. The determined cDNA sequences for these splice forms are provided in SEQ ID NO: 366-375, with the predicted amino acid sequences corresponding to the sequences of SEQ ID NO: 292, 298 and 301-303 being provided in SEQ ID NO: 299-306, respectively.

The expression levels of B305D in a variety of tumor and normal tissues were examined by real time PCR and by Northern analysis. The results indicated that B305D is highly expressed in breast tumor, prostate tumor, normal prostate tumor and normal testes, with expression being low or undetectable in all other tissues examined (colon tumor, lung tumor, ovary tumor, and normal bone marrow, colon, kidney, liver, lung, ovary, skin, small intestine, stomach).

#### EXAMPLE 12 ELICITATION OF PROSTATE TUMOR ANTIGEN-SPECIFIC CTL RESPONSES IN HUMAN BLOOD

This Example illustrates the ability of a prostate tumor antigen to elicit a CTL response in blood of normal humans.

Autologous dendritic cells (DC) were differentiated from monocyte cultures derived from PBMC of normal donors by growth for five days in RPMI medium containing 10% human serum, 50 ng/ml GMCSF and 30 ng/ml IL-4. Following culture, DC were infected overnight with recombinant P501S-expressing vaccinia virus at an M.O.I. of 5 and matured for 8 hours by the addition of 2 micrograms/ml CD40 ligand. Virus was inactivated by UV irradiation, CD8<sup>+</sup> cells were isolated by positive selection using magnetic beads, and priming cultures were initiated in 24-well plates. Following five stimulation cycles, CD8<sup>+</sup> lines were identified that specifically produced interferon-gamma when stimulated with autologous P501S-transduced fibroblasts. The P501S-specific activity of cell line 3A-1 could be maintained following additional stimulation cycles on autologous B-LCL transduced with P501S. Line 3A-1 was shown to specifically recognize autologous B-LCL transduced to

express P501S, but not EGFP-transduced autologous B-LCL, as measured by cytotoxicity assays ( $^{51}\text{Cr}$  release) and interferon-gamma production (Interferon-gamma Elispot; *see above* and Lalvani et al., *J. Exp. Med.* 186:859-865, 1997). The results of these assays are presented in Figures 6A and 6B.

### EXAMPLE 13

#### IDENTIFICATION OF PROSTATE TUMOR ANTIGENS BY MICROARRAY ANALYSIS

This Example describes the isolation of certain prostate tumor polypeptides from a prostate tumor cDNA library.

A human prostate tumor cDNA expression library as described above was screened using microarray analysis to identify clones that display at least a three fold over-expression in prostate tumor and/or normal prostate tissue, as compared to non-prostate normal tissues (not including testis). 372 clones were identified, and 319 were successfully sequenced. Table I presents a summary of these clones, which are shown in SEQ ID NOs:385-400. Of these sequences SEQ ID NOs:386, 389, 390 and 392 correspond to novel genes, and SEQ ID NOs: 393 and 396 correspond to previously identified sequences. The others (SEQ ID NOs:385, 387, 388, 391, 394, 395 and 397-400) correspond to known sequences, as shown in Table I.

Table I  
Summary of Prostate Tumor Antigens

Known Genes	Previously identified Genes	Novel Genes
T-cell gamma chain	P504S	23379 (SEQ ID NO:389)
Kallikrein	P1000C	23399 (SEQ ID NO:392)
Vector	P501S	23320 (SEQ ID NO:386)
CGI-82 protein mRNA (23319; SEQ ID NO:385)	P503S	23381 (SEQ ID NO:390)
PSA	P510S	
Ald. 6 Dehyd.	P784P	
L-Iditol-2 dehydrogenase (23376; SEQ ID NO:388)	P502S	
Ets transcription factor PDEF (22672; SEQ ID NO:398)	P706P	
hTGR (22678; SEQ ID NO:399)	19142.2, bangur.seq (22621; SEQ ID NO:396)	
KIAA0295(22685; SEQ ID NO:400)	5566.1 Wang(23404; SEQ ID NO:393)	
Prostatic Acid Phosphatase(22655; SEQ ID NO:397)	P712P	

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transglutaminase (22611; SEQ ID NO:395)	P778P	
HDLBP (23508; SEQ ID NO:394)		
CGI-69 Protein(23367; SEQ ID NO:387)		
KIAA0122(23383; SEQ ID NO:391)		
TEEG		

CGI-82 showed 4.06 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 43% of prostate tumors, 25% normal prostate, not detected in other normal tissues tested. L-iditol-2 dehydrogenase showed 4.94 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 90% of prostate tumors, 100% of normal prostate, and not detected in other normal tissues tested. Ets transcription factor PDEF showed 5.55 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 47% prostate tumors, 25% normal prostate and not detected in other normal tissues tested. hTGR1 showed 9.11 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 63% of prostate tumors and is not detected in normal tissues tested including normal prostate. KIAA0295 showed 5.59 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 47% of prostate tumors, low to undetectable in normal tissues tested including normal prostate tissues. Prostatic acid phosphatase showed 9.14 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 67% of prostate tumors, 50% of normal prostate, and not detected in other normal tissues tested. Transglutaminase showed 14.84 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 30% of prostate tumors, 50% of normal prostate, and is not detected in other normal tissues tested. High density lipoprotein binding protein (HDLBP) showed 28.06 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 97% of prostate tumors, 75% of normal prostate, and is undetectable in all other normal tissues tested. CGI-69 showed 3.56 fold over-expression in prostate tissues as compared to other normal tissues tested. It is a low abundant gene, detected in more than 90% of prostate tumors, and in 75% normal prostate tissues. The expression of this gene in normal tissues was very low. KIAA0122 showed 4.24 fold over-expression in prostate

tissues as compared to other normal tissues tested. It was over-expressed in 57% of prostate tumors, it was undetectable in all normal tissues tested including normal prostate tissues. 19142.2 bangur showed 23.25 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 97% of prostate tumors and 100% of normal prostate. It was undetectable in other normal tissues tested. 5566.1 Wang showed 3.31 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 97% of prostate tumors, 75% normal prostate and was also over-expressed in normal bone marrow, pancreas, and activated PBMC. Novel clone 23379 showed 4.86 fold over-expression in prostate tissues as compared to other normal tissues tested. It was detectable in 97% of prostate tumors and 75% normal prostate and is undetectable in all other normal tissues tested. Novel clone 23399 showed 4.09 fold over-expression in prostate tissues as compared to other normal tissues tested. It was over-expressed in 27% of prostate tumors and was undetectable in all normal tissues tested including normal prostate tissues. Novel clone 23320 showed 3.15 fold over-expression in prostate tissues as compared to other normal tissues tested. It was detectable in all prostate tumors and 50% of normal prostate tissues. It was also expressed in normal colon and trachea. Other normal tissues do not express this gene at high level.

#### EXAMPLE 14

##### IDENTIFICATION OF PROSTATE TUMOR ANTIGENS BY ELECTRONIC SUBTRACTION

This Example describes the use of an electronic subtraction technique to identify prostate tumor antigens.

Potential prostate-specific genes present in the GenBank human EST database were identified by electronic subtraction (similar to that described by Vasmatizis et al., *Proc. Natl. Acad. Sci. USA* 95:300-304, 1998). The sequences of EST clones (43,482) derived from various prostate libraries were obtained from the GenBank public human EST database. Each prostate EST sequence was used as a query sequence in a BLASTN (National Center for Biotechnology Information) search against the human EST database. All matches considered identical (length of matching sequence >100 base pairs, density of identical matches over this region > 70%) were grouped (aligned) together in a cluster. Clusters containing more than 200 ESTs were discarded since they probably represented repetitive elements or highly expressed genes such as those for ribosomal proteins. If two or more clusters shared common ESTs, those clusters were grouped together into a "supercluster," resulting in 4,345 prostate superclusters.

Records for the 479 human cDNA libraries represented in the GenBank release were downloaded to create a database of these cDNA library records. These 479 cDNA libraries were grouped into three groups, Plus (normal prostate and prostate tumor libraries, and breast cell lines, in which expression was desired), Minus (libraries from other normal adult tissues, in which expression was not desirable), and Other (fetal tissue, infant tissue, tissues found only in women, non-prostate tumors and cell lines other than prostate cell lines, in which expression was considered to be irrelevant). A summary of these library groups is presented in Table II.

**Table II**  
**Prostate cDNA Libraries and ESTs**

Library	# of Libraries	# of ESTs
Plus	25	43,482
Normal	11	18,875
Tumor	11	21,769
Cell lines	3	2,838
Minus	166	
Other	287	

Each supercluster was analyzed in terms of the ESTs within the supercluster. The tissue source of each EST clone was noted and used to classify the superclusters into four groups: Type 1- EST clones found in the Plus group libraries only; no expression detected in Minus or Other group libraries; Type 2- EST clones found in the Plus and Other group libraries only; no expression detected in the Minus group; Type 3- EST clones found in the Plus, Minus and Other group libraries, but the expression in the Plus group is higher than in either the Minus or Other groups; and Type 4- EST clones found in Plus, Minus and Other group libraries, but the expression in the Plus group is higher than the expression in the Minus group. This analysis identified 4,345 breast clusters (*see* Table III). From these clusters, 3,172 EST clones were ordered from Research Genetics, Inc., and were received as frozen glycerol stocks in 96-well plates.



**Table III**  
**Prostate Cluster Summary**

Type	# of Superclusters	# of ESTs Ordered
1	688	677
2	2899	2484
3	85	11
4	673	0
Total	4345	3172

The inserts were PCR-amplified using amino-linked PCR primers for Synteni microarray analysis. When more than one PCR product was obtained for a particular clone, that PCR product was not used for expression analysis. In total, 2,528 clones from the electronic subtraction method were analyzed by microarray analysis to identify electronic subtraction breast clones that had high tumor vs. normal tissue mRNA. Such screens were performed using a Synteni (Palo Alto, CA) microarray, according to the manufacturer's instructions (and essentially as described by Schena et al., *Proc. Natl. Acad. Sci. USA* 93:10614-10619, 1996 and Heller et al., *Proc. Natl. Acad. Sci. USA* 94:2150-2155, 1997). Within these analyses, the clones were arrayed on the chip, which was then probed with fluorescent probes generated from normal and tumor prostate cDNA, as well as various other normal tissues. The slides were scanned and the fluorescence intensity was measured.

Clones with an expression ratio greater than 3 (*i.e.*, the level in prostate tumor cDNA was at least three times the level in normal prostate cDNA) were identified as prostate tumor-specific sequences (Table IV). The sequences of these clones are provided in SEQ ID NOs:401-453, with certain novel sequences shown in SEQ ID NOs:407, 413, 416-419, 422, 426, 427 and 450.

**Table IV**  
**Prostate-tumor Specific Clones**

SEQ ID NO.	Sequence Designation	Comments
401	22545	previously identified P1000C
402	22547	previously identified P704P

403	22548	known
404	22550	known
405	22551	PSA
406	22552	prostate secretory protein 94
407	22553	novel
408	22558	previously identified P509S
409	22562	glandular kallikrein
410	22565	previously identified P1000C
411	22567	PAP
412	22568	B1006C (breast tumor antigen)
413	22570	novel
414	22571	PSA
415	22572	previously identified P706P
416	22573	novel
417	22574	novel
418	22575	novel
419	22580	novel
420	22581	PAP
421	22582	prostatic secretory protein 94
422	22583	novel
423	22584	prostatic secretory protein 94
424	22585	prostatic secretory protein 94
425	22586	known
426	22587	novel
427	22588	novel
428	22589	PAP
429	22590	known
430	22591	PSA
431	22592	known
432	22593	Previously identified P777P
433	22594	T cell receptor gamma chain
434	22595	Previously identified P705P
435	22596	Previously identified P707P
436	22847	PAP
437	22848	known
438	22849	prostatic secretory protein 57

439	22851	PAP
440	22852	PAP
441	22853	PAP
442	22854	previously identified P509S
443	22855	previously identified P705P
444	22856	previously identified P774P
445	22857	PSA
446	23601	previously identified P777P
447	23602	PSA
448	23605	PSA
449	23606	PSA
450	23612	novel
451	23614	PSA
452	23618	previously identified P1000C
453	23622	previously identified P705P

#### EXAMPLE 15

##### FURTHER IDENTIFICATION OF PROSTATE TUMOR ANTIGENS BY MICROARRAY ANALYSIS

This Example describes the isolation of additional prostate tumor polypeptides from a prostate tumor cDNA library.

A human prostate tumor cDNA expression library as described above was screened using microarray analysis to identify clones that display at least a three fold over-expression in prostate tumor and/or normal prostate tissue, as compared to non-prostate normal tissues (not including testis). 142 clones were identified and sequenced. Certain of these clones are shown in SEQ ID NOs:454-467. Of these sequences SEQ ID NOs:459-461 correspond to novel genes. The others (SEQ ID NOs:454-458 and 461-467) correspond to known sequences.

#### EXAMPLE 16

##### FURTHER CHARACTERIZATION OF PROSTATE TUMOR ANTIGEN P710P

This Example describes the full length cloning of P710P.

The prostate cDNA library described above was screened with the P710P fragment described above. One million colonies were plated on LB/Ampicillin plates. Nylon membrane filters were used to lift these colonies, and the cDNAs picked up by these filters were then denatured and cross-linked to the filters by UV light. The P710P fragment was radiolabeled and used to hybridize with the filters. Positive cDNA clones were selected and their cDNAs recovered and sequenced by an automatic ABI Sequencer. Four sequences were obtained, and are presented in SEQ ID NOs:468-471.

From the foregoing, it will be appreciated that, although specific embodiments of the invention have been described herein for the purposes of illustration, various modifications may be made without deviating from the spirit and scope of the invention. Accordingly, the present invention is not limited except as by the appended claims.

## CLAIMS

1. An isolated polypeptide comprising at least an immunogenic portion of a prostate tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472;

(b) sequences that hybridize to any of the foregoing sequences under moderately stringent conditions; and

(c) complements of any of the sequence of (a) or (b).

2. An isolated polypeptide according to claim 1, wherein the polypeptide comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472, or a complement of any of the foregoing polynucleotide sequences.

3. An isolated polypeptide comprising a sequence recited in any one of SEQ ID NO: 108, 112, 113, 114, 172, 176, 178, 327, 329, 331, 339 and 383.

4. An isolated polynucleotide encoding at least 15 amino acid residues of a prostate tumor protein, or a variant thereof that differs in one or more substitutions, deletions, additions and/or insertions such that the ability of the variant to react with antigen-specific antisera is not substantially diminished, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide comprising a sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434,

435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472, or a complement of any of the foregoing sequences.

5. An isolated polynucleotide encoding a prostate tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide comprising a sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472, or a complement of any of the foregoing sequences.

6. An isolated polynucleotide comprising a sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472.

7. An isolated polynucleotide comprising a sequence that hybridizes, under moderately stringent conditions, to a sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472.

8. An isolated polynucleotide complementary to a polynucleotide according to any one of claims 4-7.

9. An expression vector comprising a polynucleotide according to any one of claims 4-7.

10. A host cell transformed or transfected with an expression vector according to claim 9.

11. An expression vector comprising a polynucleotide according claim 8.

12. A host cell transformed or transfected with an expression vector according to claim 11.
13. A pharmaceutical composition comprising a polypeptide according to claim 1, in combination with a physiologically acceptable carrier.
14. A vaccine comprising a polypeptide according to claim 1, in combination with a non-specific immune response enhancer.
15. A vaccine according to claim 14, wherein the non-specific immune response enhancer is an adjuvant.
16. A vaccine according to claim 14, wherein the non-specific immune response enhancer induces a predominantly Type I response.
17. A pharmaceutical composition comprising a polynucleotide according to claim 4, in combination with a physiologically acceptable carrier.
18. A vaccine comprising a polynucleotide according to claim 4, in combination with a non-specific immune response enhancer.
19. A vaccine according to claim 18, wherein the non-specific immune response enhancer is an adjuvant.
20. A vaccine according to claim 18, wherein the non-specific immune response enhancer induces a predominantly Type I response.
21. An isolated antibody, or antigen-binding fragment thereof, that specifically binds to a prostate tumor protein that comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472 or a complement of any of the foregoing polynucleotide sequences.

22. A pharmaceutical composition comprising an antibody or fragment thereof according to claim 18, in combination with a physiologically acceptable carrier.

23. A pharmaceutical composition comprising an antigen-presenting cell that expresses a polypeptide according to claim 1, in combination with a pharmaceutically acceptable carrier or excipient.

24. A pharmaceutical composition according to claim 23, wherein the antigen presenting cell is a dendritic cell or a macrophage.

25. A vaccine comprising an antigen-presenting cell that expresses a polypeptide according to claim 1, in combination with a non-specific immune response enhancer.

26. A vaccine according to claim 25, wherein the non-specific immune response enhancer is an adjuvant.

27. A vaccine according to claim 25, wherein the non-specific immune response enhancer induces a predominantly Type I response.

28. A vaccine according to claim 25, wherein the antigen-presenting cell is a dendritic cell.

29. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a polypeptide according to claim 1, and thereby inhibiting the development of a cancer in the patient.

30. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a polynucleotide according to claim 4, and thereby inhibiting the development of a cancer in the patient.

31. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of an antibody or antigen-binding fragment thereof according to claim 21, and thereby inhibiting the development of a cancer in the patient.



32. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of an antigen-presenting cell that expresses a polypeptide according to claim 1, and thereby inhibiting the development of a cancer in the patient.
33. A method according to claim 32, wherein the antigen-presenting cell is a dendritic cell.
34. A method according to any one of claims 29-32, wherein the cancer is prostate cancer.
35. A fusion protein comprising at least one polypeptide according to claim 1.
36. A fusion protein according to claim 35, wherein the fusion protein comprises an expression enhancer that increases expression of the fusion protein in a host cell transfected with a polynucleotide encoding the fusion protein.
37. A fusion protein according to claim 35, wherein the fusion protein comprises a T helper epitope that is not present within the polypeptide of claim 1.
38. A fusion protein according to claim 35, wherein the fusion protein comprises an affinity tag.
39. An isolated polynucleotide encoding a fusion protein according to claim 35.
40. A pharmaceutical composition comprising a fusion protein according to claim 32, in combination with a physiologically acceptable carrier.
41. A vaccine comprising a fusion protein according to claim 35, in combination with a non-specific immune response enhancer.
42. A vaccine according to claim 41, wherein the non-specific immune response enhancer is an adjuvant.

43. A vaccine according to claim 41, wherein the non-specific immune response enhancer induces a predominantly Type I response.
44. A pharmaceutical composition comprising a polynucleotide according to claim 40, in combination with a physiologically acceptable carrier.
45. A vaccine comprising a polynucleotide according to claim 40, in combination with a non-specific immune response enhancer.
46. A vaccine according to claim 45, wherein the non-specific immune response enhancer is an adjuvant.
47. A vaccine according to claim 45, wherein the non-specific immune response enhancer induces a predominantly Type I response.
48. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a pharmaceutical composition according to claim 40 or claim 44.
49. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a vaccine according to claim 41 or claim 45.
50. A method for removing tumor cells from a biological sample, comprising contacting a biological sample with T cells that specifically react with a prostate tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:
- (i) polynucleotides recited in any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472; and
  - (ii) complements of the foregoing polynucleotides;
- wherein the step of contacting is performed under conditions and for a time sufficient to permit the removal of cells expressing the prostate tumor protein from the sample.
51. A method according to claim 50, wherein the biological sample is blood or a fraction thereof.

52. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient a biological sample treated according to the method of claim 50.

53. A method for stimulating and/or expanding T cells specific for a prostate tumor protein, comprising contacting T cells with one or more of:

- (i) a polypeptide according to claim 1;
  - (ii) a polypeptide encoded by a polynucleotide comprising a sequence provided in any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472;
  - (iii) a polynucleotide encoding a polypeptide of (i) or (ii); and/or
  - (iv) an antigen presenting cell that expresses a polypeptide of (i) or (ii);
- under conditions and for a time sufficient to permit the stimulation and/or expansion of T cells.

54. An isolated T cell population, comprising T cells prepared according to the method of claim 53.

55. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a T cell population according to claim 54.

56. A method for inhibiting the development of a cancer in a patient, comprising the steps of:

(a) incubating CD4<sup>+</sup> and/or CD8<sup>+</sup> T cells isolated from a patient with at least one component selected from the group consisting of:

- (i) a polypeptide according to claim 1;
- (ii) a polypeptide encoded by a polynucleotide comprising a sequence of any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472;
- (iii) a polynucleotide encoding a polypeptide of (i) or (ii); or
- (iv) an antigen-presenting cell that expresses a polypeptide of (i) or (ii);

such that T cells proliferate; and

(b) administering to the patient an effective amount of the proliferated T cells, and thereby inhibiting the development of a cancer in the patient.

57. A method for inhibiting the development of a cancer in a patient, comprising the steps of:

(a) incubating CD4<sup>+</sup> and/or CD8<sup>+</sup> T cells isolated from a patient with at least one component selected from the group consisting of:

- (i) a polypeptide according to claim 1;
- (ii) a polypeptide encoded by a polynucleotide comprising a sequence of any one of SEQ ID NOs: 1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472;
- (iii) a polynucleotide encoding a polypeptide of (i) or (ii); or
- (iv) an antigen-presenting cell that expresses a polypeptide of (i) or (ii);

such that T cells proliferate;

- (b) cloning at least one proliferated cell; and
- (c) administering to the patient an effective amount of the cloned T cells, and thereby inhibiting the development of a cancer in the patient.

58. A method for determining the presence or absence of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with a binding agent that binds to a prostate tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(i) polynucleotides recited in any one of SEQ ID NOs: 1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472; and

(ii) complements of the foregoing polynucleotides;

(b) detecting in the sample an amount of polypeptide that binds to the binding agent; and

(c) comparing the amount of polypeptide to a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient.

59. A method according to claim 58, wherein the binding agent is an antibody.

60. A method according to claim 59, wherein the antibody is a monoclonal antibody.

61. A method according to claim 58, wherein the cancer is prostate cancer.
62. A method for monitoring the progression of a cancer in a patient, comprising the steps of:
- (a) contacting a biological sample obtained from a patient at a first point in time with a binding agent that binds to a prostate tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472, or a complement of any of the foregoing polynucleotides;
  - (b) detecting in the sample an amount of polypeptide that binds to the binding agent;
  - (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and
  - (d) comparing the amount of polypeptide detected in step (c) to the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.
63. A method according to claim 62, wherein the binding agent is an antibody.
64. A method according to claim 63, wherein the antibody is a monoclonal antibody.
65. A method according to claim 62, wherein the cancer is a prostate cancer.
66. A method for determining the presence or absence of a cancer in a patient, comprising the steps of:
- (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a prostate tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472, or a complement of any of the foregoing polynucleotides;
  - (b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide; and

(c) comparing the amount of polynucleotide that hybridizes to the oligonucleotide to a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient.

67. A method according to claim 66, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a polymerase chain reaction.

68. A method according to claim 66, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a hybridization assay.

69. A method for monitoring the progression of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a prostate tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs:1-111, 115-171, 173-175, 177, 179-305, 307-315, 326, 328, 330, 332-335, 340-375, 381, 382 or 384-472, or a complement of any of the foregoing polynucleotides;

(b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide;

(c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and

(d) comparing the amount of polynucleotide detected in step (c) to the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

70. A method according to claim 69, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a polymerase chain reaction.

71. A method according to claim 69, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a hybridization assay.

72. A diagnostic kit, comprising:

(a) one or more antibodies according to claim 21; and

(b) a detection reagent comprising a reporter group.

73. A kit according to claim 72, wherein the antibodies are immobilized on a solid support.

74. A kit according to claim 73, wherein the solid support comprises nitrocellulose, latex or a plastic material.

75. A kit according to claim 72, wherein the detection reagent comprises an anti-immunoglobulin, protein G, protein A or lectin.

76. A kit according to claim 72, wherein the reporter group is selected from the group consisting of radioisotopes, fluorescent groups, luminescent groups, enzymes, biotin and dye particles.

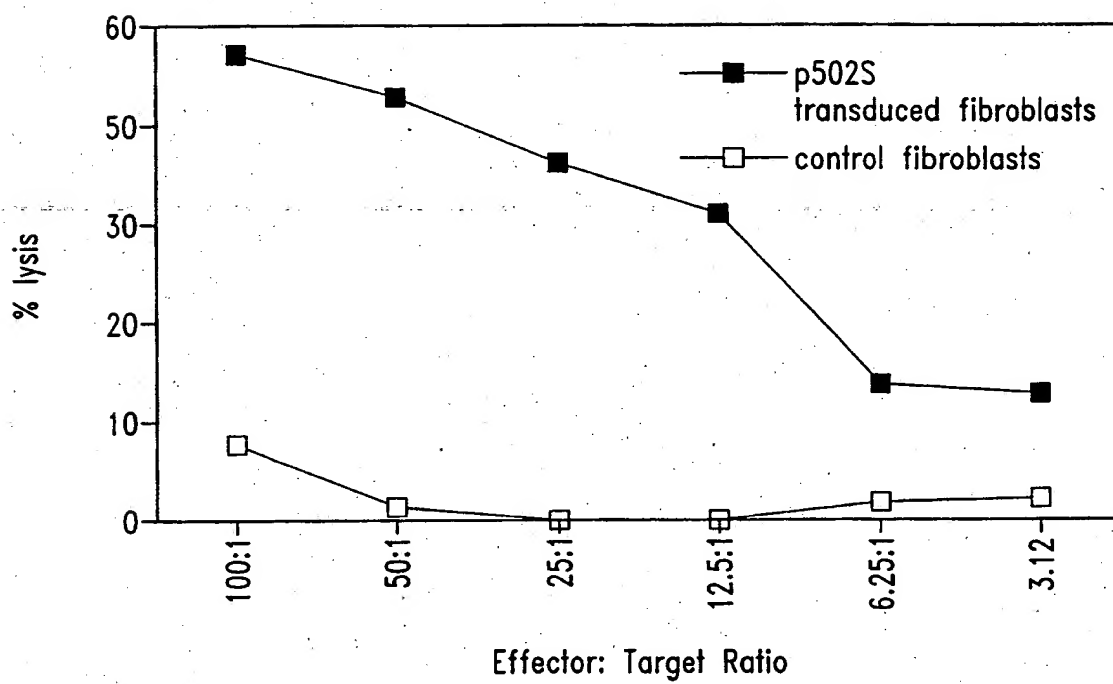
77. An oligonucleotide comprising 10 to 40 nucleotides that hybridize under moderately stringent conditions to a polynucleotide that encodes a prostate tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472, or a complement of any of the foregoing polynucleotides.

78. A oligonucleotide according to claim 77, wherein the oligonucleotide comprises 10-40 nucleotides recited in any one of SEQ ID NOs:2, 3, 8-29, 41-45, 47-52, 54-65, 70, 73-74, 79, 81, 87, 90, 92, 93, 97, 103, 104, 107, 109-111, 115-160, 171, 173-175, 177, 181, 188, 191, 193, 194, 198, 203, 204, 207, 209, 220, 222-225, 227-305, 307-315, 326, 328, 330, 332, 334, 350-365, 381, 382, 384, 386, 389, 390, 392, 393, 396, 401, 402, 407, 408, 410, 413, 415-419, 422, 426, 427, 432, 434, 435, 442-444, 446, 450, 452, 453, 459-461, 468-471 or 472.

79. A diagnostic kit, comprising:

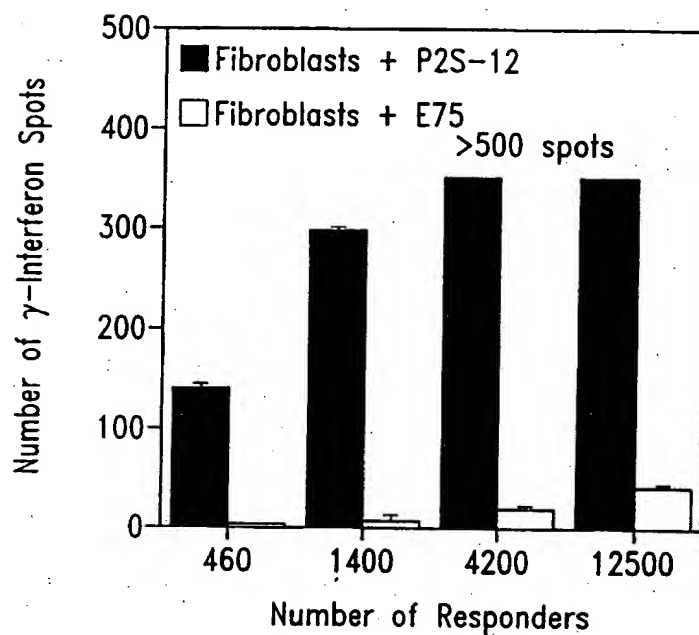
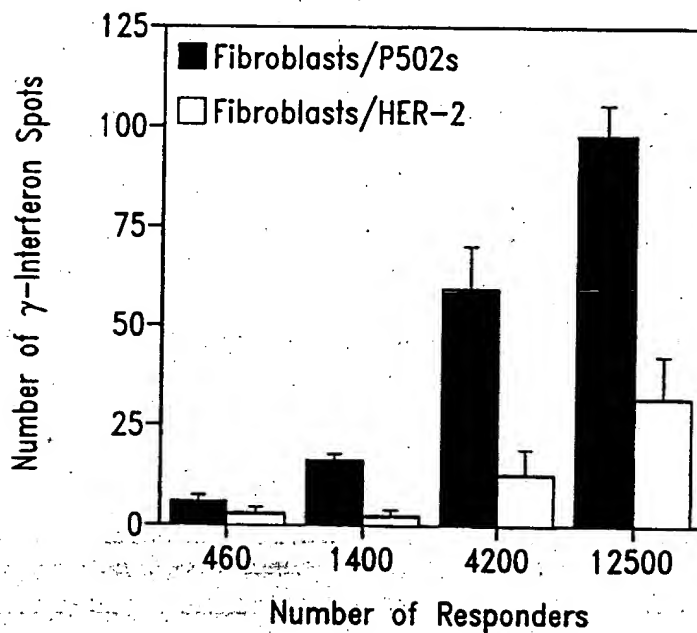
- (a) an oligonucleotide according to claim 77; and
- (b) a diagnostic reagent for use in a polymerase chain reaction or hybridization assay.

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*Fig. 1*



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*Fig. 2A**Fig. 2B*

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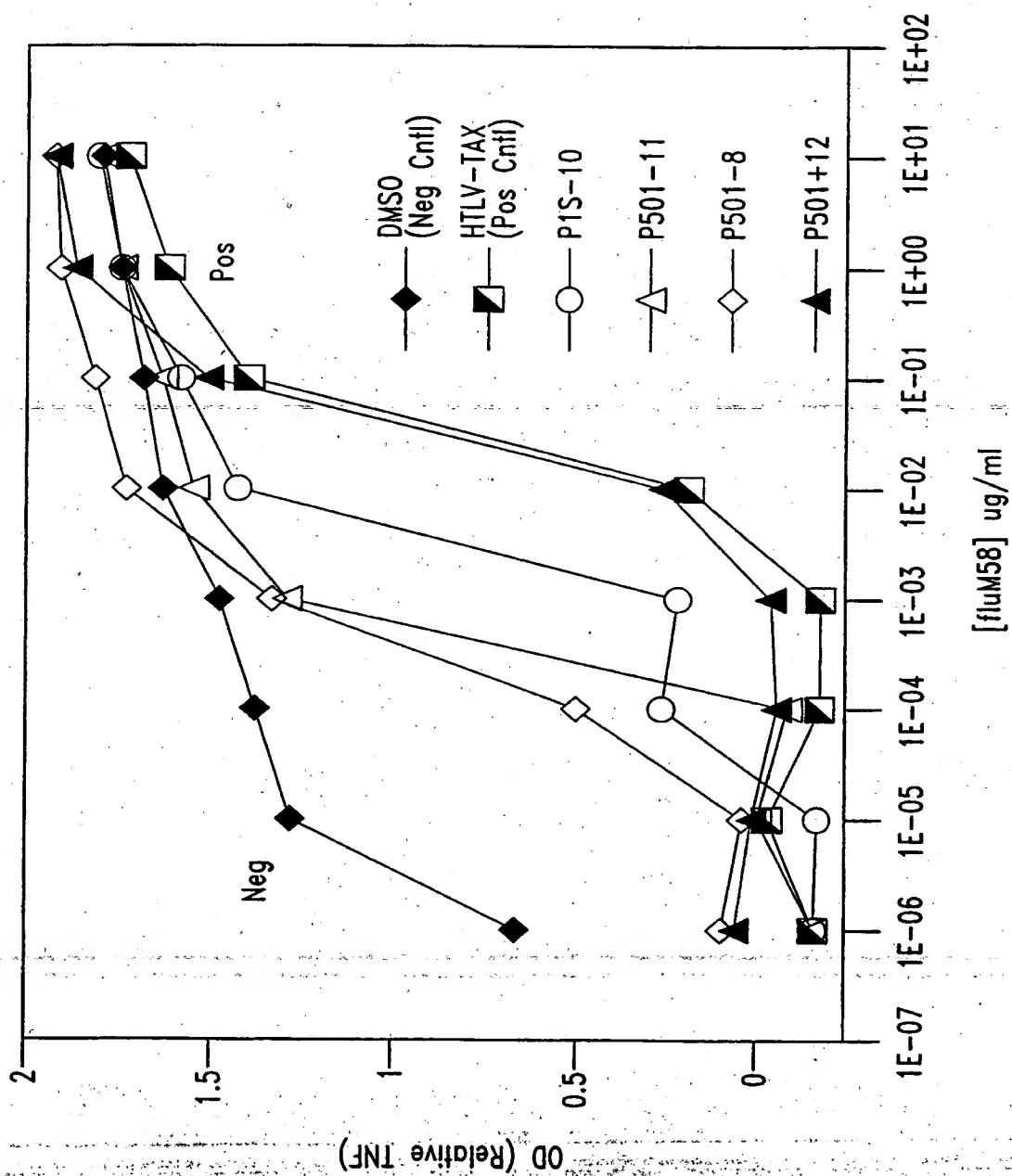
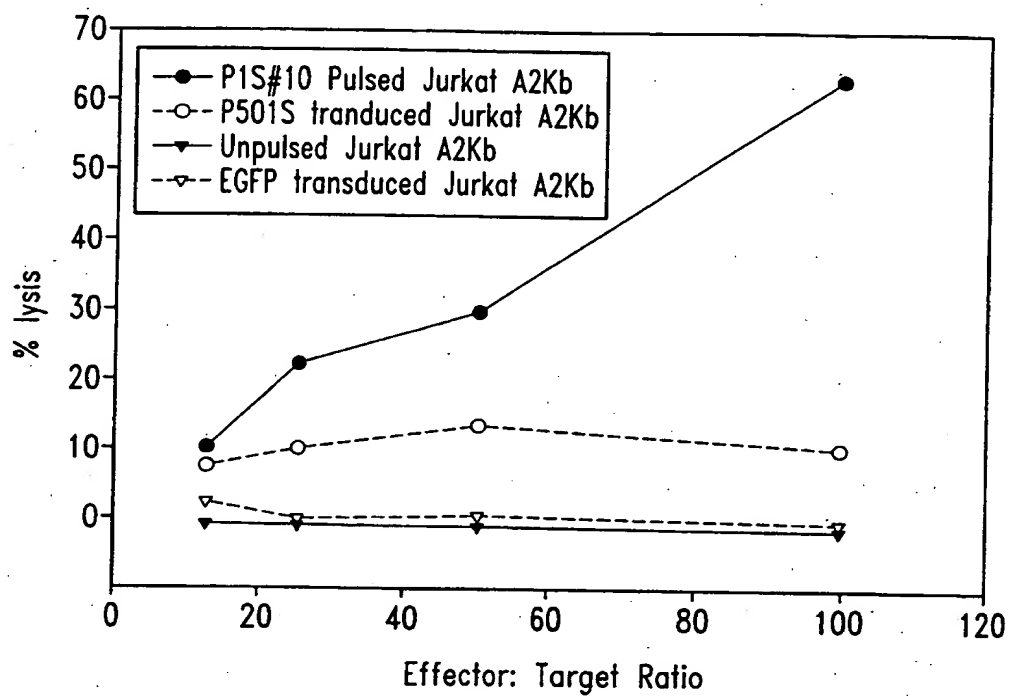
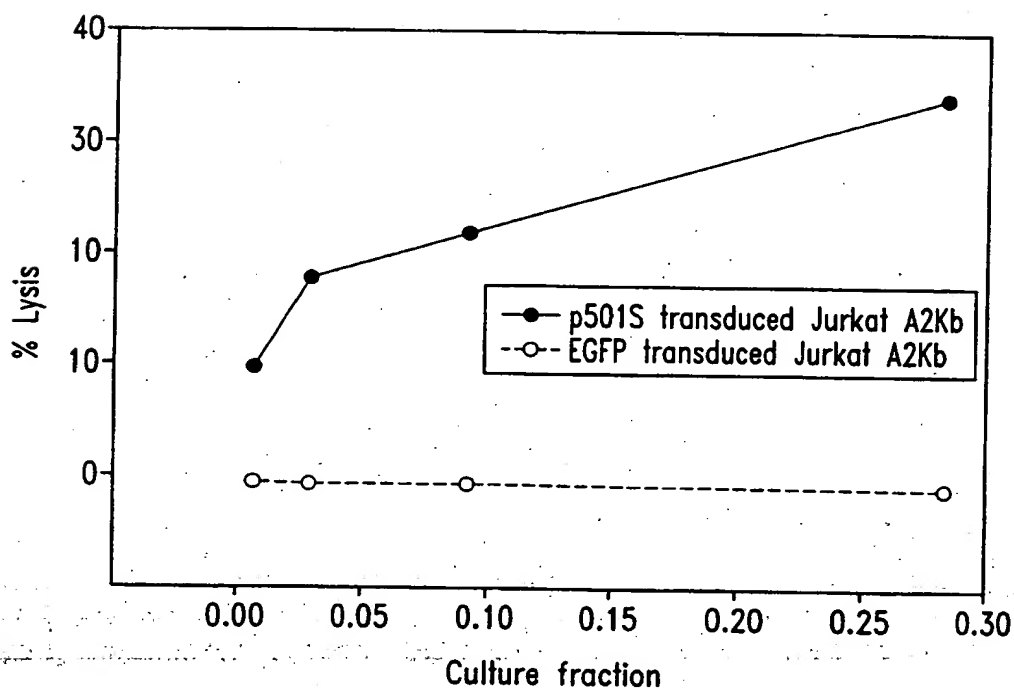


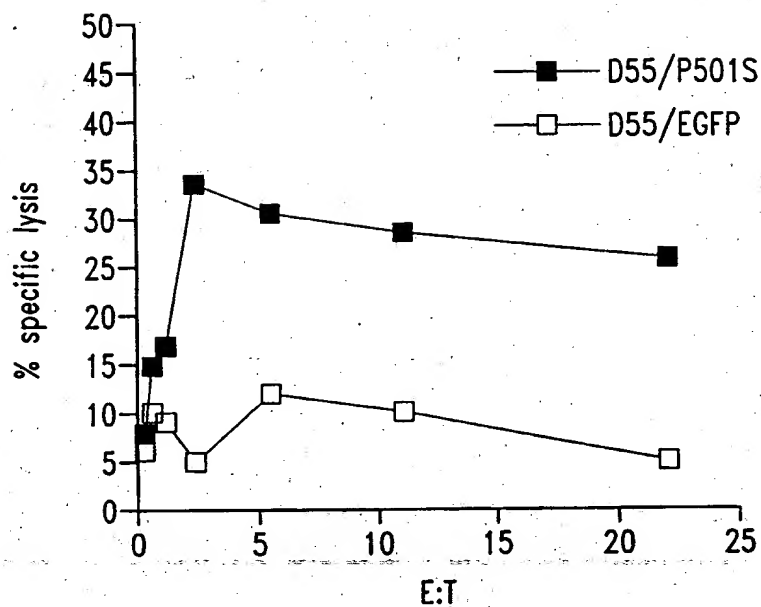
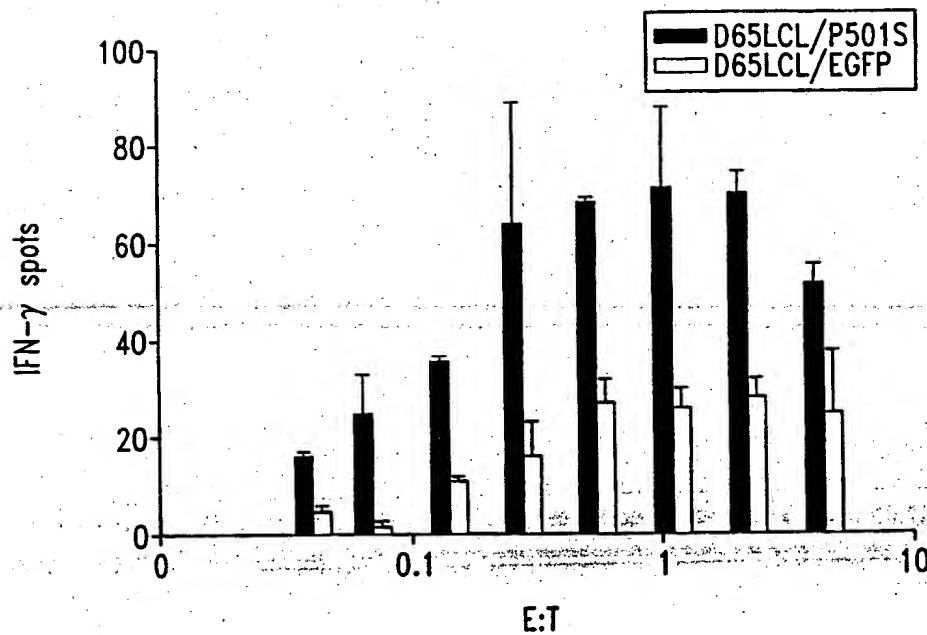
Fig. 3

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*Fig. 4**Fig. 5*

SUBSTITUTE SHEET (RULE 26)

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*Fig. 6**Fig. 7*

## SEQUENCE LISTING

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OF PROSTATE CANCER AND METHODS FOR THEIR USE

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&lt;211&gt; 773

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (773)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 3

cttttgaaag	aagggatggc	tggggtgttt	aacagcagag	gtgcagggcg	ggggctcacg	60
tcctgtcct	cactggatgat	aaacgagccc	cgttccttgt	tgtgatcatg	atgaacaacc	120
tcctcaaaag	tcagaaccgg	agtcacacag	gcattctgtg	cgtaaagat	ttgacaccac	180
tctgccttcg	tcttctttgc	aaatacatct	gcaaacttct	tcttcatttc	tgccaatca	240
tccatgctca	tctgattggg	aagttcatca	gacttttagt	canntccttt	gatcagcagc	300
tcgtagaact	ggggttctat	tgctccaaca	gccatgaatt	ccccatctgc	tgtcctgtaa	360
gtcgtataga	aagggtgctcc	accatccaac	atgttctgtc	ctcgaggggg	ggcccggtag	420
ccaattcgcc	ctatantgag	tcgtattacg	cgcgctcact	ggccgctcgt	ttacaacgtc	480
gtgactggga	aaaccctggg	cgttaccaac	ttaatcgctt	tgcagcacat	ccccctttcg	540
ccagctgggc	gtaatancca	aaaggcccgc	accgatcgcc	cttccaacag	ttgcgcacct	600
gaatgggnaa	atgggacccc	cctgttaccg	cgcattnaac	ccccgcnggg	tttngttgtt	660
acccccacnt	nnaccgctta	cactttgcc	gcgccttanc	gcccgcctcc	tttcnccttt	720
cttcccttcc	tttcncncn	ctttcccccg	gggtttcccc	cntcaaacc	cna	773

&lt;210&gt; 4

&lt;211&gt; 828

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (828)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 4

cctcctgagt	cctactgacc	tgtgctttct	ggtgtggagt	ccagggtgc	taggaaaagg	60
aatgggcaga	cacaggtgta	tgccaatgtt	tctgaaatgg	gtataatttc	gtcctctcct	120
tcggaacact	ggctgtctct	gaagacttct	cgctcagttt	cagtgaggac	acacacaaag	180
acgtgggtga	ccatgttggt	tgtgggggtgc	agagatggga	gggggtgggg	ccaccctgga	240
agagtggaca	gtgacacaag	gtggacactc	tctacagatc	actgaggata	agctggagcc	300
acaatgcatg	aggcacacac	acagcaagga	tgacnctgta	aacatagccc	acgtgtcct	360

gngggcactg	ggaagcctan	atnaggccgt	gagcanaaag	aaggggagga	tccactagtt	420
ctanagcggc	cgccaccgcg	gtgganctcc	ancctttgtt	cccttttagtg	agggttaatt	480
gcgcgcttgg	cntaatcatg	gtcatanctn	tttccctgtg	gaaattgtta	tccgctcaca	540
attccacaca	acatacganc	cggaaacata	aantgtaaac	ctgggggtgcc	taatgantga	600
ctaactcaca	ttaattgcgt	tgcgctcact	gcccgccttc	caatcnggaa	acctgtcttg	660
ccncttgcat	tnatgaatcn	gcccaacccc	ggggaaaagc	gtttgcgttt	tgggcgctct	720
tccgcttcct	cntcantta	ntccctncnc	tcggtcattc	cggctgcngc	aaaccgggtc	780
accncctcca	aaggggggtat	tccggtttcc	ccnaatccgg	gganancc		828

&lt;210&gt; 5

&lt;211&gt; 834

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(834)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 5

tttttttttt	tttttactga	tagatggaat	ttattaagct	tttcacatgt	gatagcacat	60
agttttaatt	gcatccaaag	tactaacaaa	aactctagca	atcaagaatg	gcagcatggt	120
attttataac	aatcaacacc	tgtggctttt	aaaaatttgg	tttcataaga	taattttatac	180
tgaagtaaat	ctagccatgc	ttttaaaaaa	tgctttagg	cactccaagc	ttggcagtta	240
acatttgcca	taaacaataa	taaaacaatc	acaatttaat	aaataacaaa	tacaacattg	300
taggccataa	tcatatacag	tataaggaaa	aggtggtagt	gttgagtaag	cagttattag	360
aatagaatac	cttggcctct	atgcaaatat	gtctagacac	tttgattcac	tcagccctga	420
cattcagttt	tcaaagtagg	agacagggtc	tacagtatca	ttttacagtt	tccaacacat	480
tgaaaacaag	tagaaaatga	tgagttgatt	tttattaatg	cattacatcc	tcaagagtta	540
tcaccaaccc	ctcagttata	aaaaattttc	aaagtatat	agtcataata	cttggtgtgc	600
ttatttttaa	ttagtgtctaa	atggattaag	tgaagacaac	aatgggtccc	taatgtgatt	660
gatattgggt	atttttacca	gcttctaaat	ctnaactttc	aggcttttga	actggaacat	720
tgnatnacag	tgttccanag	ttncaaccta	ctggaacatt	acagtgtgct	tgattcaaaa	780
tgttattttg	ttaaaaatta	aatttttaacc	tggtggaaaa	ataatttgaa	atna	834

&lt;210&gt; 6

&lt;211&gt; 818

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(818)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 6

tttttttttt	tttttttttt	aagaccctca	tcaatagatg	gagacataca	gaaatagtca	60
aaccacatct	acaaaatgcc	agtatcaggc	ggcggcttcg	aagccaaagt	gatgtttgga	120
tgtaaagtga	aatattagtt	ggcggatgaa	gcagatagtg	aggaaagtgt	agccaataat	180
gacgtgaagt	ccgtggaagc	ctgtggctac	aaaaaatgtt	gagccgtaga	tgccgtcgga	240
aatgggtgaag	ggagactcga	agtactctga	ggcttgtagg	agggtaaaat	agagacccag	300
taaaattgta	ataagcagtg	cttgaattat	ttggtttcgg	ttgttttcta	ttagactatg	360
gtgagctcag	gtgattgata	ctcctgatgc	gagtaatacg	gatgtgttta	ggagtgggac	420
ttctagggga	tttagcgggg	tgatgcctgt	tgggggccag	tgccctccta	gttggggggg	480
aggggctagg	ctggagtggg	aaaaggctca	gaaaaatcct	gcgaagaaaa	aaacttctga	540

ggtaataaat	aggattatcc	cgtatcgaag	gccttttttg	acaggtggtg	tgtggtggcc	600
ttggtatgtg	ctttctcgtg	ttacatcgcg	ccatcattgg	tatatgggta	gtgtgttggg	660
ttantangg	ctantatgaa	gaacttttgg	antggaatta	aatcaatngc	ttggccggaa	720
gtcattanga	nggctnaaaa	ggccctgtta	ngggctctgg	ctngggttta	cccnacccat	780
ggaatncncc	ccccggacna	ntgnatccct	attcttaa			818

&lt;210&gt; 7

&lt;211&gt; 817

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(817)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 7

tttttttttt	tttttttttt	tggctctaga	gggggtagag	ggggtgctat	agggtaaata	60
cgggccctat	ttcaaagatt	tttaggggaa	ttaattctag	gacgatgggt	atgaaactgt	120
ggtttgctcc	acagatttca	gagcattgac	cgtagtatac	ccccggtcgt	gtagcgggta	180
aagtggtttg	gttttagacgt	ccgggaattg	catctgtttt	taagcctaata	gtggggacag	240
ctcatgagtg	caagacgtct	tgtgatgtaa	ttattatacn	aatgggggct	tcaatcggga	300
gtactactcg	attgtcaacg	tcaaggagtc	gcaggtcgcc	tggttctagg	aataatgggg	360
gaagtatgta	ggaattgaag	attaatccgc	cgtagtcggt	gttctcctag	gttcaataacc	420
attggtggcc	aattgatttg	atggtaaggg	gagggatcgt	tgaactcgtc	tgttatgtaa	480
aggatncctt	ngggatggga	aggcnatnaa	ggactangga	tnaatggcgg	gcangatatt	540
tcaaacngtc	tctanttcct	gaaacgtctg	aaatgttaat	aanaattaan	tttngttatt	600
gaatnttng	gaaaagggct	tacaggacta	gaaaccaaata	angaaaanta	atnntaangg	660
cnttatcntn	aaaggtmata	accnctccta	tnatcccacc	caatngnatt	ccccacnenn	720
acnattggat	nccccanttc	canaaaanggc	cnccecccg	tgnannccnc	cttttggtcc	780
cttnantgan	ggttattcnc	ccctngcntt	atcance			817

&lt;210&gt; 8

&lt;211&gt; 799

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(799)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 8

catttccggg	ttactttct	aaggaaagcc	gagcggaagc	tgctaactgt	ggaatcggtg	60
cataaggaga	actttctgct	ggcacgcgct	agggacaagc	gggagagcga	ctccgagcgt	120
ctgaagcgca	cgtcccagaa	ggtggacttg	gcactgaaac	agctgggaca	catccgcgag	180
tacgaacagc	gcctgaaagt	gctggagcgg	gaggtccagc	agtgtagccg	cgtcctgggg	240
tgggtggcgg	angcctganc	cgtctgcct	tgctgcccc	angtgggccc	ccacccccctg	300
acctgcctgg	gtccaaacac	tgagccctgc	tggcggactt	caagganaac	ccccacangg	360
ggattttgct	cctanantaa	ggctcatctg	ggcctcgccc	ccccacctg	gttggccttg	420
tctttgagtg	gagcccatg	tccatctggg	ccactgtcng	gaccaccttt	ngggagtgtt	480
ctccttacaa	ccacannatg	ccggctcct	cccggaaacc	antcccance	tgngaaggat	540
caagnccctg	atccactnnt	nctanaaccg	gcnccnccg	cngtggaaac	cnccttntgt	600
tccttttctn	tnagggttaa	tnnccgcttg	gccttnccan	ngtccnccn	nttttccnnt	660
gttnaaattg	ttangcnccc	nccnntccn	cnnccnccan	cccgaccenn	annttnnann	720



ncctgggggt nccnnngat tgaccenncc nccctntant tgcnttnggg ncnntgccc 780  
ctttccctct nggganncg 799

<210> 9  
<211> 801  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(801)  
<223> n = A,T,C or G

<400> 9  
acgccttgat cctcccaggc tgggactggt tctgggagga gccgggcatg ctgtgggttg 60  
taangatgac actcccaaag gtggtcctga cagtggccca gatggacatg gggctcacct 120  
caaggacaag gccaccaggc gcggggggccg aagcccacat gatccttact ctatgagcaa 180  
aatccctctgt gggggcttct ccttgaagtc cgccancagg gtcagtcctt tggaccang 240  
caggatcatgg ggttgtngnc caactggggg ccncaacgca aaanggcncg gggcctcngn 300  
caccatccc angacgggc tacactnctg gacctccnc tccaccactt tcatgcgctg 360  
ttentaccg cgnatntgtc ccactgttt cngtgcenac tccancttct nggacgtgcg 420  
ctacatacgc cgggancnc nctcccgtt tgtccctatc cagtnccan caacaaattt 480  
cncctantg caccnattcc cacttttnc agntttcnc nncgngcttc cttntaaaag 540  
ggttgancgc cggaaaatnc cccaaagggg gggggccngg tacccaactn cccctnata 600  
gctgaantcc ccatnaccnn gnctcnatgg anccntcent tttaannacn ttctnaactt 660  
gggaanance ctcgncentn ccccnnttaa tcccncttg cnangnnent ccccnntcc 720  
nccnnntng gcntntnann cnaaaaaggc ccnnnancaa tctcctnnen cctcanttcg 780  
ccanccctcg aaatcgccn c 801

<210> 10  
<211> 789  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(789)  
<223> n = A,T,C or G

<400> 10  
cagtctatnt ggccagtgtg gcagctttcc ctgtggctgc cggtgccaca tgccgtgccc 60  
acagtgtggc cgtggtgaca gcttcagccg cctccaccgg gttcaccttc tcagccctgc 120  
agatcctgcc ctacacactg gcctccctct accaccggga gaagcagggt ttccctgccc 180  
aataccgagg ggacactgga ggtgctagca gtgaggacag cctgatgacc agcttcctgc 240  
caggccctaa gcctggagct cccttcctta atggacacgt ggggtgctgga ggcagtggcc 300  
tgctcccacc tccaccgcg ctctgcgggg cctctgctg tgatgtctcc gtacgtgtgg 360  
tggtgggtga gccaccgan gccagggtgg ttccgggccc gggcatctgc ctggacctcg 420  
ccatcctgga tagtgcttcc tgetgtccca ngtggcccca tccctgttta tgggctccat 480  
tgtccagctc agccagtctg tcaactgccta tatggtgtct gccgcaggcc tgggtctggg 540  
cccatttact ttgtacaca ggtantattt gacaagaacg anttggccaa atactcagcg 600  
ttaaaaaatt ccagcaacat tgggggtgga aggcctgcct cactgggtcc aactccccgc 660  
tcctgttaac cccatggggc tgccggcttg gccgccaatt tctgttgctg ccaaantnat 720  
gtggctctct gctgccacct gttgctggct gaagtgcnta cngcncanct nggggggtng 780  
gnggttccc 789

<210> 11  
 <211> 772  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(772)  
 <223> n = A,T,C or G

<400> 11

cccaccctac	ccaaatatta	gacaccaaca	cagaaaagct	agcaatggat	tcccttctac	60
tttgttaaat	aaataagtta	aatattttaa	tgcctgtgtc	tctgtgatgg	caacagaagg	120
accaacaggc	cacatcctga	taaaaggtaa	gaggggggtg	gacagcaaaa	aagacagtgc	180
tgtgggctga	ggggacctgg	ttcttgtgtg	tgccccctca	ggactcttcc	cctacaaata	240
actttcatat	gttcaaatac	catggaggag	tgtttcatcc	tagaaactcc	catgcaagag	300
ctacattaaa	cgaagctgca	ggttaagggg	cttanagatg	ggaaaccagg	tgactgagtt	360
tattcagctc	ccaaaaaccc	ttctctaggt	gtgtctcaac	taggagggta	gctgttaacc	420
ctgagcctgg	gtaatccacc	tgcagagtcc	ccgcattcca	gtgcatggaa	cccttctggc	480
ctccctgtat	aagtcagac	tgaaaccccc	ttggaaggnc	tccagtcagg	cagccctana	540
aactggggaa	aaaagaaaag	gacgccccan	ccccagctg	tgcantacg	cacctcaaca	600
gcacagggtg	gcagcaaaaa	aaccacttta	ctttggcaca	aacaaaaact	ngggggggca	660
accccgac	cccnangggg	gttaacagga	ancngggnaa	cntggaaccc	aattnaggca	720
ggccnccac	ccnaatntt	gctgggaaat	tttctctccc	ctaaattntt	tc	772

<210> 12  
 <211> 751  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(751)  
 <223> n = A,T,C or G

<400> 12

gccccaatc	cagctgccac	accacccacg	gtgactgcat	tagttcggat	gtcatacaaa	60
agctgattga	agcaaccctc	tactttttgg	tcgtgagcct	tttgcttggg	gcaggtttca	120
ttggctgtgt	tggtgacgtt	gtcattgcaa	cagaatgggg	gaaaggcact	gttctctttg	180
aagtanggtg	agtcctcaaa	atccgtatag	ttggtgaagc	cacagcactt	gagccctttc	240
atgggtggtg	tccacacttg	agtgaagtct	tcctgggaac	cataatcttt	cttgatggca	300
ggcactacca	gcaacgtcag	ggaagtgtc	agccattgtg	gtgtacacca	aggcgaccac	360
agcagctgcn	acctcagcaa	tgaagatgan	gaggangatg	aagaagaacg	tcncgagggc	420
acacttgetc	tcagtcttan	caccatanca	gccntgaaa	accaananca	aagaccacna	480
cncgggtgc	gatgaagaaa	tnacccnccg	ttgacaaact	tgcatggcac	tggganccac	540
agtggccnna	aaaatcttca	aaaaggatgc	cccactnatt	gaccccccaa	atgcccactg	600
ccaacagggg	ctgccccacn	cncnnaacga	tgancnatt	gnacaagatc	tncntggtct	660
tnatnaacnt	gaaccctgcn	tngtggtccc	tgttcaggnc	cnnngcctga	cttctnaann	720
aangaactcn	gaagncccca	cngganannc	g			751

<210> 13  
 <211> 729  
 <212> DNA  
 <213> Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(729)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 13

gagccaggcg	tccctctgcc	tgcccactca	gtggcaacac	ccgggagctg	ttttgtcctt	60
tgtggancct	cagcagtncc	ctctttcaga	actcantgcc	aaganccttg	aacaggagcc	120
accatgcagt	gcttcagctt	cattaagacc	atgatgatcc	tcttcaattt	gctcatcttt	180
ctgtgtggtg	cagccctggt	ggcagtgggc	atctgggtgt	caatcgatgg	ggcatccttt	240
ctgaagatct	tcggggccact	gtcgtccagt	gccatgcagt	ttgtcaacgt	gggctacttc	300
ctcatcgcag	ccggcggtgt	ggtcttagct	ctaggtttcc	tgggctgcta	tgggtgctaag	360
actgagagca	agtgtgccct	cgtgacgttc	ttcttcatcc	tcctcctcat	cttcattgct	420
gaggttgcaa	tgtgtgggtc	gccttggtgt	acaccacaat	ggctgagcac	ttcctgacgt	480
tgtgtgtaat	gcctgccatc	aanaaaagat	tatgggttcc	caggaanact	tactcaagt	540
ggtggaacac	caccatgaaa	gggtcaagt	gctgtggctt	cnnccaacta	tacggatttt	600
gaagantcac	ctacttcaaa	gaaaanagtg	cctttccccc	atttctgttg	caattgacaa	660
acgtccccaa	cacagccaat	tgaaaacctg	caccaaccc	aaanggggtcc	ccaaccanaa	720
attnaagg						729

&lt;210&gt; 14

&lt;211&gt; 816

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(816)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 14

tgtcttctct	caaagttggt	cttgttgcca	taacaaccac	cataggtaaa	gcgggcgcag	60
tgttcgctga	aggggttgta	gtaccagcgc	gggatgctct	ccttgcagag	tcctgtgtct	120
ggcagggtcca	cgcagtgcc	ttgtcactg	gggaaatgga	tgcgctggag	ctcgtcaaag	180
ccactcgtgt	atttttcaca	ggcagcctcg	tccgacgcgt	cggggcagtt	gggggtgtct	240
tcacactcca	ggaaactgtc	natgcagcag	ccattgctgc	agcggaaactg	ggtgggctga	300
cangtgccag	agcacactgg	atggcgccct	tccatgnnan	gggccctgng	ggaaagtccc	360
tganceccan	anctgcctct	caaangcccc	accttgccac	ccccgacagg	ctagaatgga	420
atcttcttcc	cgaaaggtag	ttnttcttgt	tgcccaancc	anccccntaa	acaaactctt	480
gcanatctgc	tccgnggggg	tcntantacc	ancgtgggaa	aagaacccca	ggcngcgaac	540
caancttggt	tggatncgaa	gcnataatct	ncnttctgc	ttggtggaca	gcaccantna	600
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gggacaagg	aantngccnt	cctttnaatt	cccnanctn	ccccctgggt	tgggggtttt	720
cncnctcta	ccccagaaan	nccgtgttcc	cccccaacta	ggggccnaaa	ccnnttnttc	780
cacaacctn	ccccacccac	gggttcngnt	ggttng			816

&lt;210&gt; 15

&lt;211&gt; 783

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(783)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 15

ccaaggcctg	ggcaggcata	nacttgaagg	tacaacccca	ggaaccctg	gtgctgaagg	60
atgtggaaaa	cacagattgg	cgctactgc	gggtgacac	ggatgtcagg	gtagagagga	120
aagacccaaa	ccaggtggaa	ctgtggggac	tcaaggaang	cacctacctg	ttccagctga	180
cagtgactag	ctcagaccac	ccagaggaca	cggccaacgt	cacagtcact	gtgctgtcca	240
ccaagcagac	agaagactac	tgcctcgcat	ccaacaangt	gggtcgctgc	cggggctctt	300
tcccacgctg	gtactatgac	cccacggagc	agatctgcaa	gagtttcgtt	tatggaggct	360
gcttgggcaa	caagaacaac	taccttcggg	aagaagagtg	cattctancc	tgtcnggggtg	420
tgcaagggtg	gcctttgana	ngcanctctg	gggctcangc	gactttcccc	cagggccctt	480
ccatggaaag	gcgccatcca	ntgttctctg	gcacctgtca	gcccaccag	ttccgctgca	540
ncaatggctg	ctgcatcnac	antttcctng	aattgtgaca	acaccccca	ntgcccccaa	600
ccctcccaac	aaagcttccc	tgttnaaaaa	tacnccantt	ggcttttnac	aaacncccg	660
cncctcctt	ttcccnntn	aacaaagggc	nctngcctt	gaactgcccn	aaccnnggaa	720
tctnccnngg	aaaaantncc	ccccctggtt	cctnnaancc	cctccncaa	anctncccc	780
ccc						783

&lt;210&gt; 16

&lt;211&gt; 801

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(801)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 16

gccccaatc	cagctgccac	accacccacg	gtgactgcat	tagttcggat	gtcatacaaa	60
agctgattga	agcaaccctc	tacttttttg	tcgtgagcct	tttgcttggt	gcagggttca	120
ttggctgtgt	tggtgacgtt	gtcattgcaa	cagaatgggg	gaaaggcact	gttctctttg	180
aagtaggggtg	agtcctcaaa	atccgtatag	ttggtgaagc	cacagcactt	gagccctttc	240
atgggtggtgt	tccacacttg	agtgaagtct	tcctgggaac	cataatcttt	cttgatggca	300
ggcactacca	gcaacgtcag	gaagtgtctc	gccattgtgg	tgtacaccaa	ggcgaccaca	360
gcagctgcaa	cctcagcaat	gaagatgagg	aggaggatga	agaagaacgt	cncgagggca	420
cacttgctct	ccgtcttagc	accatagcag	cccangaaac	caagagcaaa	gaccacaacg	480
ccngctgcca	atgaaagaaa	ntacccacgt	tgacaaaactg	catggccact	ggacgacagt	540
tggcccgaan	atcttcagaa	aagggatgcc	ccatcgattg	aacacccana	tgcccactgc	600
cnacagggct	gcncncncn	gaaagaatga	gccattgaag	aaggatcnc	ntggtcttaa	660
tgaactgaaa	ccntgcatgg	tggcccctgt	tcagggtctt	tggcagtga	ttctganaaa	720
aaggaacngc	ntnagcccc	ccaaangana	aaacaccccc	gggtgttgcc	ctgaattggc	780
ggccaaggan	ccctgccccn	g				801

&lt;210&gt; 17

&lt;211&gt; 740

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(740)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 17

gtgagagcca	ggcgtccctc	tgccctgcca	ctcagtggca	acaccggga	gctgttttgt	60
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cctttgtgga gcctcagcag ttccctcttt cagaactcac tgccaagagc cctgaacagg 120
agccaccatg cagtgtttca gcttcattaa gaccatgatg atcctcttca atttgtctcat 180
ctttctgtgt ggtgcagccc tgttggcagt gggcatctgg gtgtcaatcg atggggcatc 240
ctttctgaag atcttcgggc cactgtcgtc cagtgccatg cagtttgtca acgtgggcta 300
cttctcctc gcagccggcg ttgtggcttt tgctcttggg ttccctgggct gctatgggtgc 360
taagacggag agcaagtgtg ccctcgtgac gttcttcttc atcctcctcc tcctcttcat 420
tgctgaagtt gcagctgctg tggtcgcctt ggtgtacacc acaatggctg aaccattcct 480
gacgttgctg gtantgctg ccatcaanaa agattatggg ttcccaggaa aaattcactc 540
aantntggaa caccnccatg aaaagggtc caatttctgn tggttcccc aactataccg 600
gaattttgaa agantcnccc tacttccaaa aaaaaanant tgcttttnc cccnttctgt 660
tgcaatgaaa acntcccaan acngccaatn aaaacctgcc cnnncaaaaa ggntcncaaa 720
caaaaaaant nnaagggttn 740

```

&lt;210&gt; 18

&lt;211&gt; 802

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(802)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 18

```

ccgctgggtg cgctgggtcca gngnagccac gaagcacgtc agcatacaca gcctcaatca 60
caagggtcttc cagctgcccgc acattacgca gggcaagagc ctccagcaac actgcatatg 120
ggatacactt tacttttagca gccagggtga caactgagag gtgtcgaagc ttattcttct 180
gagcctctgt tagtggagga agattccggg cttcagctaa gtatgcagcg tatgtcccat 240
aagcaaacac tgtgagcagc cggaaggtag aggcaaatg actctcagcc agctctctaa 300
cattgggcat gtccagcagt tctccaaaca cgtagacacc agnggcctcc agcacctgat 360
ggatgagtgt ggccagcgct gcccccttgg ccgacttggc taggagcaga aattgtctct 420
ggttctgccc tgtcaccttc acttcgcgac tcatactgc actgagtgtg ggggacttgg 480
gctcaggatg tccagagacg tggttccgcc cctcncctta atgacaccgn ccanncaacc 540
gtcggtctcc gccgantgng ttcgtcgtnc ctgggtcagg gtctgctggc cncacttgc 600
aancttcgtc nggccccatg aattcacnc accggaactn gtangatcca ctnttctat 660
aaccgncgc caccgcnnt ggaactccac tctnttnc tttacttgag ggtaagggtc 720
acccttncg ttaccttggg ccaaaccntn cntgtgtcg anantngtnaa tcnggncna 780
tnccancnc atangaagcc ng 802

```

&lt;210&gt; 19

&lt;211&gt; 731

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(731)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 19

```

cnaagcttcc aggtnacggg ccgcnaanc tgaccnagg tancanaang cagnncgagg 60
gagccaccg tcacngngng gngtctttat nggagggggc ggagccacat cnetggacnt 120
cntgacccca actccccnc nncantgca gtgatgagt cagaactgaa ggtnacgtgg 180
caggaaccaa gancaaannc tgctccntc caagtcggcn nagggggcgg ggctggccac 240
gcncatccnt cnagtgtgn aaagccccnn cctgtctact tgtttgaga acngcnnga 300

```

catgcccagn	gttanataac	nggcngagag	tnantttgcc	tctcccttcc	ggctgcgcan	360
cgngtntgct	tagnggacat	aacctgacta	cttaactgaa	cccngaate	tncnccccct	420
ccactaagct	cagaacaaaa	aacttcgaca	ccactcantt	gtcacctgnc	tgctcaagta	480
aagtgtaccc	catncccaat	gtntgctnga	ngctctgncc	tgcnttangt	tcggctctgg	540
gaagacctat	caattnaagc	tatgtttctg	actgcctctt	gtcccttgn	acaancnacc	600
cnnnntcca	agggggggnc	ggcccccaat	ccccccaacc	ntnaattnan	tttanccecn	660
ccccnggcc	cggcctttta	cnancntcnn	nnacngggna	aaaccnnngc	tttncccaac	720
nnaatecncc	t					731

&lt;210&gt; 20

&lt;211&gt; 754

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(754)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 20

tttttttttt	tttttttttt	taaaaacccc	ctccattnaa	tgnaaaacttc	cgaaattgtc	60
caacccccctc	ntccaaatnn	ccntttccgg	gnggggggttc	caaacccean	ttanntttgg	120
annttaaatt	aaatnttntt	tgngggnnna	ancnaatgt	nangaaagtt	naaccanta	180
tnanttnaa	tncttgga	ccngtngntt	ccaaaaatnt	ttaaccetta	antccctccg	240
aaatngttna	nggaaaaccc	aantttctnt	aaggttggtt	gaaggntnaa	tnaaaaancc	300
nnccaattgt	ttttngccac	gcctgaatta	attggnttcc	gntgttttcc	nttaaaaana	360
ggnnancccc	ggttantnaa	tccccccnnc	cccaattata	ccganttttt	ttngaattgg	420
gancccnccg	gaattaacgg	ggnnnnntccc	tnttgggggg	cnggncccc	ccccntccgg	480
ggttngggnc	aggnncnaat	tgtttaaggg	tccgaaaaat	ccctccnaga	aaaaaanctc	540
ccaggntgag	nntnggggtt	nncccccccc	cangggccct	ctcgananagt	tgggggtttg	600
ggggcctggg	attttntttc	ccctnttncc	tcccccccc	ccnggganag	aggttngngt	660
tttgntcnn	ggccccnccn	aaganctttn	ccganttnan	ttaaatcct	gcctnggcga	720
agtccttgn	agggntaaan	ggccccctnn	cggg			754

&lt;210&gt; 21

&lt;211&gt; 755

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(755)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 21

atcancccat	gacccnaac	nggggaccnc	tcancggnc	nnnchaccnc	cgcccnatca	60
nngtagnnnc	actncnnttn	natcacnccc	cnccnactac	gccnncnanc	cnacgcnta	120
nncanatncc	actganngcg	cgangtngan	ngagaaanct	nataccanag	ncaccanacn	180
ccagctgtcc	nanaangcct	nnnatacngg	nnnatccaat	ntgnancttc	cnaagtattn	240
nncnncan	gattttcctn	anccgattac	ccntncccc	tancccttcc	cccccaacna	300
cgaaggcnct	ggncnnaagg	nngcgncc	ccgctagntc	cccnncnca	cnncncccta	360
aactcanccn	nattacnccg	ttcntgagta	tactccccg	aatctcacc	tactcaactc	420
aaaaanacn	gatacaaaat	aatncaagcc	tgnttatnac	actntgactg	ggctcttatt	480
ttagnngtcc	ntnaancntc	ctaatacttc	cagtctncct	tcnccaattt	ccnaanggct	540
ctttcngaca	gcatnttttg	gttcccnntt	gggttcttan	ngaattgccc	ttcntngaac	600

```

gggctctctt tttccttcgg ttancctggn ttcnnccggc cagttattat ttcccntttt 660
aaattctntc cntttanttt tggcnttcna aacccccggc cttgaaaacg gccccctggg 720
aaaaggttgt tttganaaaa tttttgtttt gtcc 755

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<210> 22
<211> 849
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(849)
<223> n = A,T,C or G

```

```

<400> 22
tttttttttt tttttangtg tngtcgtgca ggtagaggct tactacaant gtgaanacgt 60
acgctnggan taangcgacc cganttctag ganncnccct aaaatcanac tgtgaagatn 120
atcctgnnna cggaanggtc accggnngat nntgctaggg tgncnctcc canncnttn 180
cataactcng nggccctgcc caccaccttc ggcggcccng ngncggggcc cgggtcattn 240
gnnttaaccn cactnngcna ncggtttccn nccccnnng acccnggcga tccgggggtnc 300
tctgtcttcc cctgnagnen anaaantggg ccncggnccc ctttaccctt nnacaagcca 360
cngccttcta nccnngccc cccctccant nngggggact gccnanngt ccgttctctg 420
nnaccccnnn gggtnccctg gttgtcgant cnaccgnang ccanggatc cnaaggaagg 480
tgcgttnttg gcccctaccc ttcgctnecg nncacccttc ccgacnanga nccgctcccg 540
cncnncgnng cctcncctcg caacaccgc nctcntcngt nccggnnccc ccccaccgc 600
nccctcncnc ngncgnancn ctcnccncc gtctcannca ccacccegcc ccgccaggcc 660
ntcanccacn ggnngacnng nagnccntc gcncgcgcgn gcgnccctt cgcncngaa 720
ctnctcngg ccantnncgc tcaancenna cnaaacgccg ctgcgcggcc cgnagcgncc 780
nccctcnega gtccctccgn ctccnacc angnnttccn cgaggacacn nnaccccgcc 840
nncangcgg 849

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```

<210> 23
<211> 872
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(872)
<223> n = A,T,C or G

```

```

<400> 23
gcgcaaaacta tacttcgctc gnactcgtgc gcctcgtcnc tcttttctc cgcaaccatg 60
tctgacnanc ccgattnggc ngatatcnan aagntcganc agtccaaact gantaacaca 120
cacacnncan aganaaatcc nctgccttcc anagtanacn attgaacnng agaaccangc 180
nggcgaatcg taatnaggcg tgcgcgcgca atntgtcncc gtttatttnn ccagctcnc 240
ctnccnacc tacntctten nagctgtcnn accctngtn cgnaccccc naggtcgggg 300
tcgggtttnn nntgaccgng cnnccctcc cccntccat nacganccnc ccgcaccacc 360
nanngcncgc ncccggnct ctgcgcnc ccgtctntn cccctgtngc ctggcncngn 420
accgcattga cctcgcgnn ctncnngaaa ncnanacgt cggggttgnn annancgctg 480
tgggnngcg tctgncgcg gtcccttccn ncncttcca ccattctnt tacnggggtc 540
ccnccctc tcnncacnc cctgggacgc tntcctntgc ccccttnac tccccctt 600
cngcgtgnc cgncccccac ntcatttnca nacgntctt acaannncc ggntnnctcc 660
cnancngcn gtcancnag ggaaggngg ggnccnntg nttgacgtg ngngangtc 720
cgaanantcc tcnccntcan cnctaccct cgggcgnct ctngttnc aacttanca 780

```

ntctcccccg ngngcncttc tcagcctcnc ccnccccnct ctctgcantg tncctctgctc 840  
tnaccnntac gantnttcgn cncctctctt cc 872

<210> 24

<211> 815

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(815)

<223> n = A,T,C or G

<400> 24

gcacgcaagc	ttgagtattc	tatagngtca	cctaaatanc	ttggcntaat	catggtcnta	60
nctgncttcc	tgtgtcaaat	gtatacnaan	tanatatgaa	tctnatntga	caagannnga	120
tctnncatta	gtaacaantg	tnntgtccat	cctgtcngan	canattccca	tnnattncgn	180
cgcattcnch	gncantatn	taatngggaa	ntcnntnnn	ncaccnncat	ctatcntncc	240
gncacctgac	tggagagat	ggatnanttc	tnntntgacc	nacatgttca	tcttggattn	300
aanancccc	cgcngnccac	cggttngnng	cnagccnntc	ccaagacctc	ctgtggaggt	360
aacctgcgtc	aganncatca	aacntgggaa	acccgcnncc	angtnnaagt	ngnnncanan	420
gatcccgacc	agnttnnacc	atcccttcnc	agcgcacctc	ttngtgcctt	anagnngnagc	480
gtgtccnanc	cnetcaacat	ganacgcgcc	agnccanccg	caattnggca	caatgtcgnc	540
gaaccccccta	gggggantna	tncaaanccc	caggattgtc	cncncangaa	atcccnanc	600
ccnccccctac	ccncttttgg	gacngtgacc	aantccccga	gtcccagtc	ggccngnctc	660
ccccaccggt	nnccttgggg	gggtgaanct	cngnntcanc	cngnccgagg	ntcgnaagga	720
accggncctn	ggncgaanng	ancnntcnga	agnccnctnt	cgtataacce	cccccncca	780
nccnacngnt	agntcccccc	cngggtnccg	aangg			815

<210> 25

<211> 775

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(775)

<223> n = A,T,C or G

<400> 25

ccgagatgtc	tcgctccgtg	gccttagctg	tgctcgcgct	actctctctt	tctggcctgg	60
aggctatcca	gcgtactcca	aagattcagg	tttactcacg	tcattccagca	gagaatggaa	120
agtcaaat	cctgaattgc	tatgtgtctg	ggtttcatcc	atccgacatt	gaanttgtact	180
tactgaagaa	tgganagaga	attgaaaaag	tggagcattc	agacttgtct	tccagcaagg	240
actggtcttt	ctatctcntg	tactacactg	aattcacccc	cactgaaaaa	gatgagtatg	300
cctgccgtgt	gaaccatgtg	actttgtcac	agcccaagat	agttaagtgg	gatcgagaca	360
tgtaagcagn	cnnatgga	gtttgaagat	gccgcatttg	gattggatga	attccaaatt	420
ctgcttgctt	gcntttta	antgatatgc	ntatacaccc	taccctttat	gnccccaaat	480
tgtaggggtt	acatnantgt	tcnctntngga	catgatcttc	ctttataant	ccnccnttcg	540
aattgcccgt	cncnngttn	ngaagtgttc	cnaaccacg	gttggctccc	ccaggtcncc	600
tcttacggaa	gggcctgggc	cnccttncaa	ggttggggga	accnaaaatt	tcnctnttgc	660
ccncccncca	cnntcttng	nncncanttt	ggaacccttc	cnattcccc	tggcctenna	720
nccttnncta	anaaaacttn	aaancgtngc	naaanntttt	acttcccccc	ttacc	775

<210> 26



<211> 820  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(820)  
 <223> n = A,T,C or G

<400> 26

anattantac	agtgtaatct	tttcccagag	gtgtgtanag	ggaacggggc	ctagaggcat	60
cccanagata	ncttatanca	acagtgcctt	gaccaagagc	tgctgggcac	atttcctgca	120
gaaaagggtg	cgggtcccat	cactcctcct	ctcccatagc	catcccagag	gggtgagtag	180
ccatcangcc	ttcgggtggga	gggagtcang	gaaacaacan	accacagagc	anacagacca	240
ntgatgacca	tgggcgggag	cgagcctcct	ccctgnaccg	gggtggcana	nganagccta	300
nctgaggggt	cacactataa	acgttaacga	ccnagatnan	cacctgcctc	aagtgcaccc	360
ttcctacctg	acnaccagng	accnnaact	gcngcctggg	gacagcctg	ggancagcta	420
acnnagcact	cacctgcccc	cccattggcg	tnegentccc	tggtcctgnc	aagggaagct	480
ccctgttgga	attncgggga	naccaaggga	ncctcctcct	ccanctgtga	aggaaaaann	540
gatggaattt	tncccttccg	gcnntcccc	tcttcttcta	cacgccccct	nntactctc	600
tccctctntt	ntcctgnenc	acttttnacc	ccnnnatttc	ccttnattga	tcggannctn	660
ganattccac	tnncgectnc	cntcnatcng	naanacnaaa	nactntctna	ccnnggggat	720
gggnncctcg	ntcatcctct	cttttctnct	accnccnntt	ctttgcctct	ccttngatca	
780tccaacntc	gntggccntn	ccccccnnn	tcttttnecc			

820

<210> 27  
 <211> 818  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(818)  
 <223> n = A,T,C or G

<400> 27

tctgggtgat	ggcctcttcc	tcctcagggg	cctctgactg	ctctggggcca	aagaatctct	60
tgtttcttct	ccgagcccca	ggcagcgggt	attcagccct	gccccacctg	attctgatga	120
ctgcggatgc	tgtgacggac	ccaaggggca	aatagggtcc	caggggtccag	ggagggggcg	180
ctgctgagca	cttcgcggcc	tcaccctgcc	cagccctg	catgagctct	gggctgggtc	240
tccgcctcca	gggttctgct	cttccangca	ngccancaag	tggcgctggg	ccacactggc	300
ttcttctg	ccctcctg	gctctganc	tctgtcttcc	tgctctgtgc	angcnccttg	360
gatctcagtt	tcctcctctc	anngaactct	gtttctgann	tcttcantta	actntgant	420
tatnaccnan	tggnetgtnc	tgctcnactt	taatgggcn	gaccggctaa	tcctccttc	480
ntcccttcc	anttcnnna	accngcttnc	cntctctcc	ccntancecg	ccngggaanc	540
ctcctttgcc	ctnaccangg	gcnnnnaccg	ccctnnctn	ggggggcnng	gtnnctnnc	600
ctgntnnccc	cnetcnent	tnctctgtcc	cnnnnnccg	nngcannttc	ncngtcccn	660
tnnctcttcn	ngntcgnaa	ngntcnentn	tnnnnnnncn	ngntnnntcn	tcctctcnc	720
cnnntgnang	tnnttnnnnc	ncngnncccc	nnnnnnnnnn	nggnntnnnn	tcctcncngc	780
ccnncccc	ngnattaagg	cctccnntct	ccggcenc			818

<210> 28  
 <211> 731  
 <212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(731)

<223> n = A,T,C or G

<400> 28

aggaagggcg	gagggatatt	gtangggatt	gagggatagg	agnataangg	gggaggtgtg	60
tccaacatg	anggtgnngt	tctcttttga	angaggggtg	ngtttttann	ccnggtgggt	120
gattnaaccc	cattgtatgg	agnnaaagg	tttnagggat	ttttcggctc	ttatcagtat	180
ntanattcct	gtnaatcgga	aaatnatntt	tcnnnnggaa	aatnttgctc	ccatccgnaa	240
attnctcccc	ggtagtgcg	nttngggggg	cngccangtt	tcccaggctg	ctanaatcgt	300
actaaagntt	naagtgggan	tncaaataaa	aacctnnac	agagnatccn	tacccgactg	360
tnnnttncct	tcgcccctng	actctgcng	agcccaatac	ccnngngnat	gtcncccngn	420
nnngcgnnc	tgaaannnnc	tcngggctnn	gancatcang	gggtttcgca	tcaaaagcnn	480
cgttttncat	naaggcactt	tngcctcatc	caaccnctng	ccctcnncca	tttngccgctc	540
nggttncct	acgctnntng	cncctnnntn	ganattttnc	ccgcttnggg	naancctect	600
gnaatgggta	gggncttntc	ttttnacnnc	gnggtntact	aatcnnctnc	acgctntnctt	660
tctcnacccc	cccccttttt	caatcccanc	ggcnaatggg	gtctccccnn	cgangggggg	720
nnccccannc	c					731

<210> 29

<211> 822

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(822)

<223> n = A,T,C or G

<400> 29

actagtccag	tgtgggtggaa	ttccattgtg	ttggggncnc	ttctatgant	antnttagat	60
cgctcanacc	tcacancctc	ccnacnangc	ctataangaa	nannaataga	nctgtncnnt	120
atntntacnc	tcatanncct	cnnnaccac	tcctctttaa	ccctactgt	gcctatngcn	180
tnnctantct	ntgcgcctn	cnaaccacn	gtgggcnac	cncnngnat	ctcnatctcc	240
tcnccatntn	gcctananta	ngtncatacc	ctatacctac	nccaatgcta	nnnctaancn	300
tccatnantt	annntaacta	ccactgaant	ngactttcnc	atnancctct	aatttgaatc	360
tactctgact	cccacngcct	annnattagc	ancntcccc	nacnatntct	caaccaaate	420
ntcaacaacc	tatctanctg	ttcnccaacc	nttnccctcg	atccccnnac	aacccccctc	480
ccaaataccc	nccacctgac	ncctaaccn	caccatcccc	gcaagccnan	ggncatttan	540
ccactggaat	cacnatngga	naaaaaaac	ccnaactctc	tancncnnat	ctccctaana	600
aatnctcctn	naatttactn	ncantnccat	caancccaac	tgaaacnnaa	ccccgttttt	660
tanatccctt	ctttcgaaaa	ccnacccttt	annncccaac	ctttnggggc	ccccnctnc	720
ccnaatgaag	gncnccaat	cnangaaacg	ncntgaaaa	ancnaggcna	anannntccg	780
canatcctat	cccttanttn	ggggnccctt	nccnggggcc	cc		822

<210> 30

<211> 787

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

&lt;222&gt; (1) ... (787)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 30

cgccgcctg	ctctggcaca	tgccctcctga	atggcatcaa	aagtgatgga	ctgcccattg	60
ctagagaaga	ccttctctcc	tactgtcatt	atggagccct	gcagactgag	ggctcccctt	120
gtctgcagga	tttgatgtct	gaagtcgtgg	agtgtggctt	ggagctcctc	atctacatna	180
gctggaagcc	ctggagggcc	tctctcgcca	gcctccccct	tctctccacg	ctctccangg	240
acaccagggg	ctccaggcag	cccattatc	ccagnangac	atggtgtttc	tccacgcgga	300
cccattgggg	ctgnaaggcc	agggtctcct	ttgacacccat	ctctcccgtc	ctgcctggca	360
ggccgtggga	tccactantt	ctanaacggn	cgccaccncg	gtgggagctc	cagcttttgt	420
tcccnttaat	gaagggttaat	tgcnegcttg	gcgtaatcat	nggtcanaac	tntttcctgt	480
gtgaaattgt	ttntccccctc	ncnattccnc	ncnacatacn	aaccgcgaan	cataaagtgt	540
taaagcctgg	gggtngcctn	nngaanaac	tnaactcaat	taattgcgtt	ggctcatggc	600
ccgctttccn	ttcnggaaaa	ctgtctctcc	ctgcnttntt	gaatcgcca	ccccccnggg	660
aaaagcgggt	tgcnttttng	gggntcctt	ccntctcccc	cctcnctaen	ccctncgctt	720
cggtcgttnc	nggtngcggg	gaangggnat	nnctccccnc	naagggggng	agnnngntat	780
ccccaaa						787

&lt;210&gt; 31

&lt;211&gt; 799

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (799)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 31

tttttttttt	tttttttggc	gatgctactg	tttaattgca	ggaggtgggg	gtgtgtgtac	60
catgtaccag	ggctattaga	agcaagaagg	aaggagggag	ggcagagcgc	cctgctgagc	120
aacaaaggac	tcctgcagcc	ttctctgtct	gtctcttggc	gcaggcacat	ggggaggcct	180
cccgagggtt	gggggccacc	agtccagggg	tgggagcact	acanggggtg	ggagtggggtg	240
gtggctggtt	cnaatggcct	gncacanatc	cctacgatc	ttgacacctg	gatttcacca	300
ggggaccttc	tgttctccca	nggnaacttc	ntnnatctcn	aaagaacaca	actgtttctt	360
cngcanttct	ggctgttcat	ggaaagcaca	ggtgtccnat	ttnggctggg	acttggtaca	420
tatggttccg	gcccacctct	ccentcnaen	aagtaattca	ccccccccc	ccntctnttg	480
cctgggccct	taantaccca	caccgggaact	canttantta	ttcatcttng	gntgggcttg	540
ntnatcncn	cctgaangcg	caaagttgaa	aggccacgcc	gtncnccnctc	cccatagnan	600
nttttntnt	canctaattg	ccccccnggc	aacnatccaa	tcccccccn	tgggggcccc	660
agcccanggc	ccccgntctg	ggnnnccngn	cncgnantcc	ccaggntctc	ccantcngnc	720
ccnnngcnc	cccgcacgca	gaacanaagg	ntngagccnc	cgcannnnnn	nggtnnnac	780
ctcgcccccc	ccnnegnnng					799

&lt;210&gt; 32

&lt;211&gt; 789

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (789)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 32

tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	60
ttttnccnag	ggcagggtta	ttgacaacct	cncgggacac	aancaggctg	gggacaggac	120
ggcaacaggc	tccggcggcg	gcggcggcgg	ccctacctgc	ggtaccaa	ntgcagcctc	180
cgctcccgt	tgatnttct	ctgcagctgc	aggatgccnt	aaaacagggc	ctcgccntn	240
ggtgggcacc	ctgggatttn	aatttccacg	ggcacaatgc	ggtcgcancc	cctcaccacc	300
nattaggaat	agtggtnnta	ccnccnccg	ttggcncact	ccccntggaa	accacttntc	360
gcggctccgg	catctggtct	taaaccttgc	aaacnctggg	gccctctttt	tggttantnt	420
nccngccaca	atcatnactc	agactggcnc	gggctggccc	caaaaaancn	ccccaaaacc	480
ggnccatgtc	ttnnccgggt	tgctgcnatn	tncatcacct	cccgggcnca	ncaggncaac	540
ccaaaagttc	ttngggcccn	caaaaaanct	ccggggggnc	ccagtttcaa	caaagtcac	600
ccccctggcc	cccaaactct	ccccccgntt	nctgggtttg	ggaaccacg	cctctnnctt	660
tggnnggcaa	gntggntccc	ccttcggggc	cccggtgggc	ccnnctctaa	ngaaaacncc	720
ntcctnnnca	ccatcccccc	nngnnacgnc	tancaangna	tccctttttt	tanaaacggg	780
ccccccnccg						789

&lt;210&gt; 33

&lt;211&gt; 793

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (793)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 33

gacagaacat	gttggatggt	ggagcacctt	tctatacgac	ttacaggaca	gcagatgggg	60
aattcatggc	tggttgagca	atanaacccc	agttctacga	gctgctgac	aaaggacttg	120
gactaaagtc	tgatgaactt	cccaatcaga	tgagcatgga	tgattggcca	gaaatgaana	180
agaagtttgc	agatgtattt	gcaaagaaga	cgaaggcaga	gtggtgtcaa	atctttgacg	240
gcacagatgc	ctgtgtgact	ccggttctga	cttttgagga	ggttggtcat	catgatcaca	300
acaangaacg	gggctcgttt	atcaccantg	aggagcagga	cgtgagcccc	cgccctgcac	360
ctctgctgtt	aaacacccca	gccatccctt	ctttcaaaag	ggatccacta	cttctagagc	420
ggncgccacc	gcgggtggagc	tccagctttt	gttcccttta	gtgagggtta	attgcgcgct	480
tggcgtaatc	atggtcatan	ctgtttcctg	tgtgaaattg	ttatccgctc	acaattccac	540
acaacatacg	anccggaagc	atnaaatttt	aaagcctggn	ggtngcctaa	tgantgaact	600
nactcacatt	aattggcttt	gcgctcactg	cccgttttcc	agtccggaaa	acctgtcctt	660
gccagctgcc	nttaatgaat	cnggccaccc	cccggggaaa	aggcngtttg	cttnttgggg	720
cgcncttccc	gctttctcgc	ttcctgaant	ccttcccccc	ggtctttcgg	cttgcggcna	780
acggtatcna	cct					793

&lt;210&gt; 34

&lt;211&gt; 756

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (756)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 34

gccgcgaccg	gcatgtacga	gcaactcaag	ggcgagtggga	accgtaaaaag	ccccaatctt	60
ancaagtgcg	gggaanagct	gggtcgactc	aagctagttc	ttctggagct	caacttcttg	120

ccaaccacag	ggaccaagct	gaccaaacag	cagctaattc	tggcccgtga	catactggag	180
atcgggggccc	aatggagcat	cctacgcaan	gacatccct	ccttcgagcg	ctacatggcc	240
cagctcaa	gctactactt	tgattacaan	gagcagctcc	ccgagtcagc	ctatatgcac	300
cagctcttgg	gcctcaacct	cctcttcctg	ctgtcccaga	accgggtggc	tgantnccac	360
acgganttgg	ancggctgcc	tgcccaanga	catacanacc	aatgtctaca	tcnaccacca	420
gtgtcctgga	gcaatactga	tgganggcag	ctaccncaa	gtnttcctgg	ccnagggtaa	480
catccccgc	cgagagctac	accttcttca	ttgacatcct	gctcgacact	atcaggggatg	540
aaaatcgcn	ggttgctcca	gaaaggctnc	aanaanatcc	tttctnctga	aggcccccg	600
atnctnctagt	nctagaatcg	gcccgcctac	gcggtgganc	ctccaacctt	tcgttncct	660
ttactgaggg	ttnattgccg	cccttggcgt	tatcatggtc	acnccngttn	cctgtgttga	720
aattnttaac	ccccacaa	tccacgccna	cattng			756

&lt;210&gt; 35

&lt;211&gt; 834

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(834)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 35

ggggatctct	anactnacct	gnatgcatgg	ttgtcggtgt	ggtcgctgtc	gatgaanatg	60
aacaggatct	tgcccttgaa	gctctcggt	gctgtnttta	agttgctcag	tctgccgtca	120
tagtcagaca	cncctcttggg	caaaaaacan	caggatntga	gtcttgattt	cacctccaat	180
aatcttcngg	gctgtctgct	cggtgaactc	gatgaanang	ggcagctgg	tgtgtntgat	240
aaantccanc	angttctcct	tggtgacctc	cccttcaaag	ttgttcggc	cttcatcaaa	300
cttctnnaan	angannancc	canctttgtc	gagctggnat	ttgganaaca	cgtcaccgtt	360
ggaaactgat	cccaaattgg	atgtcatcca	tcgctctgc	tgcccgcaaa	aaacttgctt	420
ggcncaaate	cgactcccn	tccttgaaag	aagccnatca	cacccccctc	cctggactcc	480
nncaangact	ctnccgctnc	ccntccng	cagggttgg	ggcannccgg	gccntgccc	540
ttcttcagcc	agttcaacat	nttcatcagc	ccctctgcca	gctgtntat	tccttggggg	600
ggaanccgct	tctcccttcc	tgaannaact	ttgaccgtng	gaatagccgc	gntcnccnt	660
acntnctggg	ccgggttcaa	antccctcen	ttgncntcn	cctcgggcca	ttctggattt	720
nccnaacttt	ttcttcccc	cncccnccg	ngtttgntt	tttcatnggg	ccccaaactt	780
gctnttggcc	antcccttgg	gggcntntan	cnccctnt	ggtcctntng	ggcc	834

&lt;210&gt; 36

&lt;211&gt; 814

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(814)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 36

cgngcgttt	cnngccgcgc	ccggtttcca	tgacnaaggc	tccttcang	ttaaatacn	60
cctagnaaac	attaatgggt	tgctctacta	atacatcata	cnaaccagta	agcctgcccc	120
naacgccaac	tcaggccatt	cctaccaaag	gaagaaaggc	tggtctctcc	acccctgtga	180
ggaaaggcct	gccttgtaag	acaccacaat	ncggctgaat	ctnaagtctt	gtgttttact	240
aatggaaaaa	aaaaataaac	aanaggtttt	gttctcatgg	ctgccaccgc	cagcctggca	300
ctaaaacanc	ccagcgctca	cttctgcttg	ganaaatatt	ctttgctctt	ttggacatca	360

ggcttgatgg	tatcaactgcc	acntttccac	ccagctgggc	ncccttcccc	catntttgtc	420
antganctgg	aaggcctgaa	ncttagtctc	caaaagtctc	ngcccacaag	accggccacc	480
aggggagntc	ntttncagtg	gatctgccaa	anantaccn	tatcatcnnt	gaataaaaag	540
gccccgaac	ganatgcttc	cancancctt	taagacccat	aatcctngaa	ccatgggtgcc	600
cttcgggtct	gacccnaaag	gaatgttctt	gggtcccant	ccctcctttg	ttntctacgt	660
tgtnttggac	ccntgctngn	atnacccaan	tganatcccc	ngaagcacc	tnccctggc	720
atttganttt	cntaaattct	ctgccctaen	nctgaaagca	cnattccctn	ggcncnaan	780
ggngaactca	agaaggtctn	ngaaaaacca	cncn			814

&lt;210&gt; 37

&lt;211&gt; 760

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(760)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 37

gcattgctgct	cttcctcaaa	gttggtcttg	ttgccataac	aaccaccata	ggtaaagcgg	60
gcgcagtgtt	cgctgaaggg	gtttagtagt	cagcgcgagg	tgctctcctt	gcagagtcct	120
gtgtctggca	ggccacgca	atgccctttg	tactggggga	aatggatgcg	ctggagctcg	180
tcnaanccac	tcgtgtattt	ttcacangca	gcctcctccg	aagcctccgg	gcagttgggg	240
gtgtcgtcac	actccactaa	actgtcgatn	cancagccca	ttgctgcagc	ggaactgggt	300
gggctgacag	gtgccagaac	acactggatn	ggcctttcca	tggaagggcc	tgggggaaat	360
cncctnancc	caaactgcct	ctcaaaggcc	accttgcaac	ccccgacagg	ctagaaatgc	420
actcttcttc	ccaaaggtag	ttgttcttgt	tgcccaagca	ncctccanca	aaccaaaanc	480
ttgcaaaatc	tgctccgtgg	gggtcatnnn	taccanggtt	ggggaaanaa	accggcngn	540
ganccncctt	gtttgaatgc	naaggnaata	atcctcctgt	cttgcttggg	tggaaanagca	600
caattgaact	gttaacnttg	ggcnggttc	cncnnggtg	gtctgaaact	aatcacgcgc	660
actggaaaaa	ggtangtgcc	ttccttgaat	tcccaaannt	ccccngnttt	tgggtntttt	720
ctcctctncc	ctaaaaatcg	tnttcccccc	ccntangggc			760

&lt;210&gt; 38

&lt;211&gt; 724

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(724)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 38

tttttttttt	tttttttttt	tttttttttt	tttttaaaaa	ccccctccat	tgaatgaaaa	60
cttcnnaaat	tgtccaaccc	cctcnnccaa	atnnccattt	ccgggggggg	gttccaaacc	120
caaattaatt	ttgganttta	aattaaatnt	tnattngggg	aanaanccaa	atgtnaagaa	180
aatttaaccc	attatnaact	taaatncctn	gaaacccntg	gnttccaaaa	atttttaacc	240
cttaaatccc	tccgaaattg	ntaanggaaa	accaaattcn	cctaaggctn	tttgaaggtt	300
ngattttaac	ccccttnant	tnttttnacc	cnnngctnaa	ntatttngnt	tccgggtgtt	360
tcctnttaan	cntnggtaac	tcccngntat	gaannncctt	aanccaatta	aaccgaattt	420
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cccnctttcg	gggtttgggn	ntaggttgaa	tttttnnang	ncccaaaaaa	ncccccaana	540
aaaaaactcc	caagnnttaa	ttngaantnc	ccccctccca	ggccttttgg	gaaaggnggg	600

tttntggggg ccngggantt cnttcccccn ttncnccccc cccccnggt aaanggttat	660
ngnntttggt ttttgggccc cttnanggac cttccggatn gaaattaaat ccccggnecg	720
gccg	724

&lt;210&gt; 39

&lt;211&gt; 751

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(751)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 39

tttttttttt tttttctttg ctcacattta atttttatatt tgattttttt taatgctgca	60
caacacaata tttatttcat ttgtttcttt tatttcattt tatttgtttg ctgctgctgt	120
tttattttatt tttactgaaa gtgagaggga actttttgtgg ccttttttcc tttttctgta	180
ggccgcctta agctttctaa atttggaaca tctaagcaag ctgaanggaa aaggggggtt	240
cgcaaaatca ctccgggggaa nggaaagggt gctttgttaa tcatgcccta tgggtgggtga	300
ttaactgctt gtacaattac ntttcacttt taattaattg tgctnaangc ttaattana	360
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tcccggcnnt cnttgaaaca cacngcngaa ngttctcatt ntccccncnc caggtnaaaa	480
tgaagggtta ccatntttta cncacacctc acntggcnnn gcctgaatcc tcnaaaaancn	540
ccttcaancn aattnctnng ccccggtcnc gcntnngtc cnccegggt cggggaantn	600
cacccccnga anncnntnnc naacnaaatt ccgaaaatat tcccnntcnc tcaattcccc	660
cnnagactnt cctcnncnnc cncaattttt tttnttcac gaacncgnnc cnaaaaatgn	720
nnnnncctc cnetngtccn naatcnccan c	751

&lt;210&gt; 40

&lt;211&gt; 753

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(753)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 40

gtggtatttt ctgtaagatc aggtgttccct ccctcgtagg tttagaggaa acaccctcat	60
agatgaaaac cccccgaga cagcagcact gcaactgcca agcagccggg gtaggagggg	120
cgccctatgc acagctgggc ccttgagaca gcagggttc gatgtcaggc tcgatgtcaa	180
tggctctggaa gcggcggtg tacctgcgta ggggcacacc gtcagggcc accaggaact	240
tctcaaagt ccaggcaacn tcgttgcgac acaccggaga ccaggatn agcttgggt	300
cggtcataan cgcggtggcg tcgtcgttg gagctggcag ggcctccgc aggaaggcna	360
ataaaagggt cgccccgca cgttcact cgcacttctc naanaccatg angttgggt	420
cnaaccacc accannccgg acttccttga nggaattccc aaatctcttc gntcttgggc	480
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aaanccaccn cctcctctt tcatctgggt tntntcccc ggacctgggt tctctcaag	600
ggancccata tctnaccan tactcactt nccccccnt gnnaccanc cttctanngn	660
ttccncccc ncctctggcc cntcaaan gcttnacna cctgggtctg ccttcccccc	720
tnccctatct gnacccnncn tttgtctcan tnt	753

&lt;210&gt; 41

<211> 341  
 <212> DNA  
 <213> Homo sapien

<400> 41  
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 ttctttaaac cttgttcatt atgaacactg aaaataggaa tttgtgaaga gttaaaaagt 180  
 tatagcttgt ttacgtagta agtttttgaa gtctacattc aatccagaca cttagttagag 240  
 tgttaaactg tgatttttaa aaaatatcat ttgagaatat tctttcagag gtattttcat 300  
 ttttactttt tgattaattg tgttttatat attagggtag t 341

<210> 42  
 <211> 101  
 <212> DNA  
 <213> Homo sapien

<400> 42  
 acttactgaa tttagtctg tgctcttctt tatttagtgt tgtatcataa atactttgat 60  
 gtttcaaaca ttctaaataa ataattttca gtggcttcat a 101

<210> 43  
 <211> 305  
 <212> DNA  
 <213> Homo sapien

<400> 43  
 acatctttgt tacagtctaa gatgtgttct taaatcacca tctcttctg gtctcaccc 60  
 tccaggtggt tctcacactg taattagagc tattgaggag tctttacagc aaattaagat 120  
 tcagatgctt tgctaagtct agagttctag agttatgttt cagaaagtct aagaaaccca 180  
 cctcttgaga ggtcagtaaa gaggacttaa tatttcatat ctacaaaatg accacaggat 240  
 tggtacaga acgagagtta tcttgataa ctacagagct agtacctgcc cgggggccgc 300  
 tcgaa 305

<210> 44  
 <211> 852  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (852)  
 <223> n = A,T,C or G

<400> 44  
 acataaatat cagagaaaag tagtctttga aatatttacg tccaggagtt ctttgtttct 60  
 gattatttgg tgtgtgtttt ggtttgtgtc caaagtattg gcagcttcag ttttcatttt 120  
 ctctccatcc tcgggcattc ttcccaaatt tatataccag tcttcgtcca tccacacgct 180  
 ccagaatttc tctttttagt taatatctca tagctcggct gagcttttca taggtcatgc 240  
 tgctgttgtt cttcttttta cccatagct gagccactgc ctctgatttc aagaacctga 300  
 agacgccctc agatcggctt tccattttta ttaatcctgg gttcttgtct gggttcaaga 360  
 ggatgtcgcg gatgaattcc cataagttag tccctctcgg gttgtgcttt ttgggtgtggc 420  
 acttggcagg ggggtcttgc tcttttttca tatcaggtga ctctgcaaca ggaaggtgac 480  
 tgggtggtgt catggagatc tgagccgggc agaaagtttt gctgtccaac aaatctactg 540  
 tgctaccata gttggtgtca tataaatagt tctngtcttt ccagggtgttc atgatggaag 600



gctcagtttg	ttcagtcttg	acaatgacat	tgtgtgtgga	ctggaacagg	tcactactgc	660
actggccggt	ccacttcaga	tgctgcaagt	tgctgtagag	gagntgcccc	gccgtccctg	720
ccgcccgggt	gaactcctgc	aaactcatgc	tgcaaagggtg	ctcgccgttg	atgtcgaact	780
cntggaaagg	gatacaattg	gcatccagct	ggttgggtgtc	caggagggtga	tggagccact	840
cccacacctg	gt					852

&lt;210&gt; 45

&lt;211&gt; 234

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 45

acaacagacc	cttgctcgct	aacgacctca	tgctcatcaa	gttggacgaa	tccgtgtccg	60
agtctgacac	catccggagc	atcagcattg	cttcgcagtg	ccctaccgcg	gggaactctt	120
gcctcgtttc	tggctggggg	ctgctggcga	acggcagaat	gcctaccgtg	ctgcagtgcg	180
tgaacgtgtc	gggtggtgtct	gaggagggtct	gcagtaagct	ctatgacccg	ctgt	234

&lt;210&gt; 46

&lt;211&gt; 590

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (590)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 46

actttttatt	taaatgttta	taaggcagat	ctatgagaat	gatagaaaac	atgggtgtgta	60
atttgatagc	aatatttttg	agattacaga	gttttagtaa	ttaccaatta	cacagttaaa	120
aagaagataa	tatattccaa	gcanatacaa	aatatcta	gaaagatcaa	ggcaggaaaa	180
tgantataac	taattgacaa	tggaaaatca	attttaatgt	gaattgcaca	ttatccttta	240
aaagctttca	aaanaanaa	ttattgcagt	ctanttaatt	caaacagtgt	taaatgggtat	300
caggataaan	aactgaaggg	canaaaagaat	taattttcac	ttcatgtaac	ncacccanat	360
ttacaatggc	ttaaattgcan	ggaaaaagca	gtggaagtag	ggaagtantc	aagggtctttc	420
tggctctctaa	tctgccttac	tctttgggtg	tggctttgat	cctctggaga	cagctgccag	480
ggctcctgtt	atatccacaa	tcccagcagc	aagatgaagg	gatgaaaaag	gacacatgct	540
gccttccttt	gaggagactt	catctcactg	gccaacactc	agtcacatgt		590

&lt;210&gt; 47

&lt;211&gt; 774

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (774)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 47

acaagggggc	ataatgaagg	agtggggana	gattttaaag	aaggaaaaaa	aacgaggccc	60
tgaacagaat	tttctgnac	aacggggcct	caaaataatt	ttcttgggga	ggttcaagac	120
gcttcactgc	ttgaaactta	aatggatgtg	ggacanaatt	ttctgtaatg	accctgaggg	180
cattacagac	gggactctgg	gaggaaggat	aaacagaaag	gggacaaagg	ctaatacccaa	240
aacatcaaag	aaaggaaggt	ggcgtcatac	ctcccagcct	acacagttct	ccagggtctct	300

```

cctcatccct ggaggacgac agtggaggaa caactgacca tgtccccagg ctctgtgtg 360
ctggctcctg gtcttcagcc cccagctctg gaagcccacc ctctgctgat cctgcgtggc 420
ccacactcct tgaacacaca tccccagggt atattcctgg acatggctga acctcctatt 480
cctacttccg agatgccttg ctccctgcag cctgtcaaaa tcccaactac cctccaaacc 540
acggcatggg aagcctttct gacttgcttg attactccag catcttgga caatccctga 600
ttccccactc cttagaggca agataggggt gttaagagta gggctggacc acttgagacc 660
aggctgctgg cttcaaattn tggctcattt acgagctatg ggaccttggg caagtnatct 720
tcacttctat gggcntcatt ttgttctacc tgcaaaatgg gggataataa tagt 774

```

```

<210> 48
<211> 124
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(124)
<223> n = A,T,C or G

```

```

<400> 48
canaaattga aattttataa aaaggcattt ttctcttata tccataaaat gatataattt 60
ttgcaantat anaaatgtgt cataaattat aatgttcctt aattacagct caacgcaact 120
tggt 124

```

```

<210> 49
<211> 147
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(147)
<223> n = A,T,C or G

```

```

<400> 49
gccgatgcta ctattttatt gcaggagggt ggggtgtttt tattattctc tcaacagctt 60
tgtggctaca ggtgggtgtct gactgcatna aaaanttttt tacgggtgat tgcaaaaatt 120
ttagggcacc catatcccaa gcantgt 147

```

```

<210> 50
<211> 107
<212> DNA
<213> Homo sapien

```

```

<400> 50
acattaaatt aataaaagga ctgttgggggt tctgctaaaa cacatggctt gatataattgc 60
atgggttgag gttaggagga gttaggcata tgttttggga gaggggt 107

```

```

<210> 51
<211> 204
<212> DNA
<213> Homo sapien

```

```

<400> 51
gtcctaggaa gtctagggga cacacgactc tggggtcacg gggccgacac acttgcacgg 60

```

cggggaaggaa aggcagagaa gtgacaccgt caggggggaaa tgacagaaag gaaaatcaag 120  
gccttgcaag gtcagaaagg ggactcaggg cttccaccac agccctgccc cacttggcc 180  
cctccctttt gggaccagca atgt 204

<210> 52

<211> 491

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(491)

<223> n = A,T,C or G

<400> 52

acaaagataa catatatctt ataacaaaaa tttgatagtt ttaaagggtta gtattgtgta 60  
gggtattttt caaaagacta aagagataac tcaggtaaaa agttagaaat gtataaaaca 120  
ccatcagaca gggtttttaa aaacaacata ttacaaaatt agacaatcat ccttataaaaa 180  
aaaacttctt gtatcaattt cttttgttca aaatgactga cttaantatt tttaaatatt 240  
tcanaaacac ttcctcaaaa attttcaana tggtagcttt canatgtnc ctcagtccca 300  
atgttgctca gataaataaa tctcgtgaga acttaccacc caccacaagc tttctggggc 360  
atgcaacagt gtcttttctt tnccttttct tttttttttt ttacaggcac agaaactcat 420  
caattttatt tggataacaa aggggtctcca aattatattg aaaaataaat ccaagttaat 480  
atcactcttg t 491

<210> 53

<211> 484

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(484)

<223> n = A,T,C or G

<400> 53

acataattta gcagggctaa ttaccataag atgctattta ttaanaggtn tatgatctga 60  
gtattaacag ttgctgaagt ttggtatttt tatgcagcat tttctttttg ctttgataac 120  
actacagaac ccttaaggac actgaaaatt agtaagtaaa gttcagaaac attagctgct 180  
caatcaaact tctacataac actatagtaa ttaaaacggt aaaaaaaagt gttgaaatct 240  
gcactagtat anaccgctcc tgtcaggata anactgcttt ggaacagaaa gggaaaaanc 300  
agcttttgant ttctttgtgc tgatangagg aaaggctgaa ttaccttggt gcctctccct 360  
aatgattggc aggtcnggta aatnccaaaa catattccaa ctcaacactt cttttccnng 420  
tancttgant ctgtgtattc caggancagg cggatggaat gggccagccc ncggatgttc 480  
cant 484

<210> 54

<211> 151

<212> DNA

<213> Homo sapien

<400> 54

actaaacctc gtgcttgatga actccatata gaaaacgggt ccatccctga acacggctgg 60  
ccactgggta tactgctgac aaccgcaaca aaaaaaacac aaatccttgg cactggctag 120  
tctatgtcct ctcaagtgcc tttttgtttg t 151

<210> 55  
<211> 91  
<212> DNA  
<213> Homo sapien

<400> 55  
acctggcttg tctccgggtg gttcccggcg cccccacgg tccccagaac ggacactttc 60  
gccctccagt ggatactcga gccaaagtgg t 91

<210> 56  
<211> 133  
<212> DNA  
<213> Homo sapien

<400> 56  
ggcggatgtg cgttggttat atacaaatat gtcattttat gtaagggact tgagtatact 60  
tggttttttg gtatctgtgg gttgggggga cgtccagga accaataccc catggatacc 120  
aagggacaac tgt 133

<210> 57  
<211> 147  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(147)  
<223> n = A,T,C or G

<400> 57  
actctggaga acctgagccg ctgctccgcc tctgggatga ggtgatgcan gcngtggcgc 60  
gactgggagc tgagcccttc cctttgcgcc tgcctcagag gattgttgcc gacntgcana 120  
tctcantggg ctggatncat gcagggt 147

<210> 58  
<211> 198  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(198)  
<223> n = A,T,C or G

<400> 58  
acagggatat aggtttnaag ttattgtnat tgtaaaatac attgaatttt ctgtatactc 60  
tgattacata catttatcct taaaaaaga tgtaaatctt aatttttatg ccatctatta 120  
attaccaat gagttacctt gtaaatgaga agtcatgata gcactgaatt ttaactagtt 180  
ttgacttcta agtttggg 198

<210> 59  
<211> 330  
<212> DNA  
<213> Homo sapien

&lt;400&gt; 59

acaacaaatg gggtgtgagg aagtcttatac agcaaaaactg gtgatggcta ctgaaaagat	60
ccattgaaaa ttatcattaa tgatttttaa tgacaagtta tcaaaaactc actcaatttt	120
cacctgtgct agcttgctaa aatgggagtt aactctagag caaatatagt atcttctgaa	180
tacagtcaat aaatgacaaa gccagggcct acaggtgggt tccagacttt ccagaccag	240
cagaaggaat ctattttatc acatggatct ccgtctgtgc tcaaaaatacc taatgatatt	300
tttcgtcttt attggacttc tttgaagagt	330

&lt;210&gt; 60

&lt;211&gt; 175

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 60

accgtgggtg cttctacat tcttgacggc tcttcacca acatctgggt ctacttcggc	60
gtcgtgggtc cttctctct cactctcatc cagctgggtc tgctcatcga ctttgcgac	120
tcttgaacc agcgggtggc gggcaaggcc gaggagtgcg attcccgtgc ctggt	175

&lt;210&gt; 61

&lt;211&gt; 154

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 61

accccacttt tctcctgtg agcagctctgg acttctcact gctacatgat gagggtgagt	60
gggtgttgct cttcaacagt atcctccct tccggatct gctgagccgg acagcagtcg	120
tggactgcac agccccgggg ctccacattg ctgt	154

&lt;210&gt; 62

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 62

cgtcagagcc ctatagtgag tcgtattaga	30
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&lt;210&gt; 63

&lt;211&gt; 89

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 63

acaagtcatt tcagcacctc ttgctcttca aaactgacca tcttttatat ttaatgcttc	60
ctgtatgaat aaaaatgggt atgtcaagt	89

&lt;210&gt; 64

&lt;211&gt; 97

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 64

accggagtaa ctgagtcggg acgctgaatc tgaatccacc aataaataaa gggtctgcag	60
aatcagtgca tccaggattg gtccttgat ctggggt	97

<210> 65  
 <211> 377  
 <212> DNA  
 <213> Homo sapien  
  
 <220>  
 <221> misc\_feature  
 <222> (1)...(377)  
 <223> n = A,T,C or G

<400> 65  
 acaacaanaa ntcccttctt taggccactg atggaaacct ggaacccccct tttgatggca 60  
 gcatggcgctc ctaggccttg acacagcggc tgggggtttgg gctntcccaa accgcacacc 120  
 ccaaccctgg tctaccaca nttctggcta tgggctgtct ctgccactga acatcagggt 180  
 tcggtcataa natgaaatcc caanggggac agaggctcagt agaggaagct caatgagaaa 240  
 ggtgctgttt gctcagccag aaaacagctg cctggcattc gccgctgaac tatgaacccg 300  
 tgggggtgaa ctaccccccag gaggaatcat gcctgggcga tgcaanggtg ccaacaggag 360  
 gggcgggagg agcatgt 377

<210> 66  
 <211> 305  
 <212> DNA  
 <213> Homo sapien

<400> 66  
 acgcctttcc ctcagaattc agggaagaga ctgtgcctg ccttcctccg ttgttgcgctg 60  
 agaaccctgt tgccttcc caccatattc accctcgctc catctttgaa ctcaaacacg 120  
 aggaactaac tgcaccctgg tctctcccc agtccccagt tcaccctcca tccctcacct 180  
 tctccactc taagggatat caacactgcc cagcacaggg gccctgaatt tatgtggttt 240  
 ttatatattt tttaataaga tgcactttat gtcatttttt aataaagtct gaagaattac 300  
 tgttt 305

<210> 67  
 <211> 385  
 <212> DNA  
 <213> Homo sapien

<400> 67  
 actacacaca ctccacttgc cttgtgaga cactttgtcc cagcacttta ggaatgctga 60  
 ggtcggacca gccacatctc atgtgcaaga ttgccagca gacatcaggt ctgagagttc 120  
 ccctttttaa aaaggggact tgcttaaaaa agaagtctag ccacgattgt gtgagcagc 180  
 tgtgctgtgc tggagattca cttttgagag agttctctc tgagacctga tctttagagg 240  
 ctgggcagtc ttgcacatga gatggggctg gtctgatctc agcactcctt agtctgcttg 300  
 cctctcccag ggccccagcc tggccacacc tgcttacagg gcactctcag atgccatac 360  
 catagtttct gtgctagtgg accgt 385

<210> 68  
 <211> 73  
 <212> DNA  
 <213> Homo sapien

<400> 68  
 acttaaccag atatattttt accccagatg gggatattct ttgtaaaaaa tgaaaataaa 60  
 gtttttttaa tgg 73

<210> 69  
 <211> 536  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(536)  
 <223> n = A,T,C or G

<400> 69  
 actagtccag tgtggtggaa ttccattgtg ttggggggtc tcaccctcct ctctgcagc 60  
 tccagctttg tgctctgcct ctgaggagac catggcccag catctgagta ccctgctgct 120  
 cctgctggcc accctagctg tggccctggc ctggagcccc aaggaggagg ataggataat 180  
 cccgggtggc atctataacg cagacctcaa tgatgagtgg gtacagcgtg cccttcactt 240  
 cgccatcagc gagtataaca aggccaccaa agatgactac tacagacgtc cgctgcgggt 300  
 actaagagcc aggaacaga ccgttggggg ggtgaattac ttcttcgacg tagagggtgg 360  
 ccgaaccata tgtaccaagt ccagcccaa cttggacacc tgtgccttcc atgaacagcc 420  
 agaactgcag aagaacagt tgtgctcttt cgagatctac gaagtccct ggggagaaca 480  
 gaangtcctt gggtgaaatc caggtgtcaa gaaatcctan ggatctgttg ccaggc 536

<210> 70  
 <211> 477  
 <212> DNA  
 <213> Homo sapien

<400> 70  
 atgacccta acagggggccc tctcagccct cctaatagacc tccggcctag ccatgtgatt 60  
 tcacttccac tccataacgc tccatcactt aggcctacta accaacacac taaccatata 120  
 ccaatgatgg cgcgatgtaa cagagaaaag cacataccaa ggccaccaca caccacctgt 180  
 ccaaaaaggc cttcgatacg ggataatcct atttattacc tcagaagttt ttttcttcgc 240  
 agggattttt ctgagccttt taccactcca gcctagcccc taccceccaa ctaggaggggc 300  
 actggcccc aacaggcatc accccgctaa atcccctaga agtcccactc ctaaacacat 360  
 ccgtattact cgcacagga gtatcaatca cctgagctca ccatagtcta atagaaaaca 420  
 accgaaacca aattattcaa agcactgctt attacaattt tactgggtct ctatttt 477

<210> 71  
 <211> 533  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(533)  
 <223> n = A,T,C or G

<400> 71  
 agagctatag gtacagtgtg atctcagctt tgcaaacaca ttttctacat agatagtact 60  
 aggtattaat agatatgtaa agaaagaaat cacaccatta ataatggtaa gattgggtta 120  
 tgtgatttta gtggtatttt tggcaccctt atatatgttt tccaaacttt cagcagtgat 180  
 attatttcca taacttaaaa agtgagtttg aaaaagaaaa tctccagcaa gcattctcatt 240  
 taaataaagg tttgtcatct ttaaaaatac agcaatatgt gactttttta aaaagctgtc 300  
 aaatagggtg gaccctacta ataattatta gaaatacatt taaaaacatc gagtacctca 360  
 agtcagtttg ccttgaaaaa tatcaaatat aactcttaga gaaatgtaca taaaagaatg 420  
 cttcgtaatt ttggagtang aggttccttc ctcaattttg tattttttaa aagtacatgg 480  
 taaaaaaaaa aattcacaaac agtatataag gctgtaaaat gaagaattct gcc 533

<210> 72  
 <211> 511  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(511)  
 <223> n = A,T,C or G

<400> 72  
 tattacggaa aaacacacca cataattcaa ctancaaaga anactgcttc agggcgtgta 60  
 aaatgaaagg cttccaggca gttatctgat taaagaacac taaaagaggg acaaggctaa 120  
 aagccgcagg atgtctacac tatancaggc gctatttggt ttggctggag gagctgtgga 180  
 aaacatggan agattgggtgc tgganatcgc cgtggctatt ctcattggtt attacanagt 240  
 gaggttctct gtgtgcccac tggtttgaaa accgttctnc aataatgata gaatagtaca 300  
 cacatgagaa ctgaaatggc ccaaacccag aaagaaagcc caactagatc ctcagaanac 360  
 gcttctaggg acaataaccg atgaagaaaa gatggcctcc ttgtgcccc gtctgttatg 420  
 atttctctcc attgcagcna naaacccgtt cttctaagca aacncagggt atgatggcna 480  
 aaatacaccc cctcttgaag naccnggagg a 511

<210> 73  
 <211> 499  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(499)  
 <223> n = A,T,C or G

<400> 73  
 cagtgccagc actggtgcca gtaccagtag caataacagt gccagtgccg gtgccagcac 60  
 cagtgggtggc ttcagtgtctg gtgccagcct gaccgccact ctcacatttg ggctcttcgc 120  
 tggccttggg ggagctgggt ccagcaccag tggcagctct ggtgcctgtg gtttctccta 180  
 caagttagat tttagatatt gttaatcctg ccagctcttc tcttcaagcc aggggtgcac 240  
 ctcagaaacc tactcaacac agcactctag gcagccacta tcaatcaatt gaagttgaca 300  
 ctctgcatta aatctatttg ccatttctga aaaaaaaaaa aaaaaaaggc cggccgctcg 360  
 antctagagg gcccggttaa acccgctgat cagcctcgac tgtgccttct anttgccagc 420  
 catctgttgt ttgccccctc cccgntgcct tccttgacct tggaaagtgc cactcccact 480  
 gtcctttcct aantaaat 499

<210> 74  
 <211> 537  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(537)  
 <223> n = A,T,C or G

<400> 74  
 tttcatagga gaacacactg aggagatact tgaagaattt ggattcagcc gcgaagagat 60



ttatcagctt	aactcagata	aaatcattga	aagtaataag	gtaaaagcta	gtctctaact	120
tccaggccca	cggctcaagt	gaatttgaat	actgcattta	cagtgtagag	taacacataa	180
cattgtatgc	atggaaacat	ggaggaacag	tattacagtg	tcctaccact	ctaatcaaga	240
aaagaattac	agactctgat	tctacagtga	tgattgaatt	ctaaaaatgg	taatcattag	300
ggcttttgat	ttataanact	ttgggtactt	atactaaatt	atggtagtta	tactgccttc	360
cagtttgctt	gatataattt	ttgatattaa	gattcttgac	ttatattttg	aatgggttct	420
actgaaaaan	gaatgatata	ttcttgaaga	catcgatata	catttattta	cactcttgat	480
tctacaatgt	agaaaatgaa	ggaaatgccc	caaattgtat	gggtataaaa	gtcccgt	537

&lt;210&gt; 75

&lt;211&gt; 467

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(467)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 75

caaanacaat	tgttcaaaag	atgcaaata	tacactactg	ctgcagctca	caaacacctc	60
tgcatattac	acgtacctcc	tcctgctcct	caagtagtgt	ggctctat	gccatcatca	120
cctgctgtct	gcttagaaga	acggctttct	gctgcaangg	agagaaatca	taacagacgg	180
tggcacaagg	aggccatctt	ttcctcatcg	gttattgtcc	ctagaagcgt	cttctgagga	240
tctagttggg	ctttctttct	gggtttgggc	catttcantt	ctcatgtgtg	tactattcta	300
tcattattgt	ataacggttt	tcaaaccngt	gggcacncag	agaacctcac	tctgtaataa	360
caatgaggaa	tagccacggg	gatctccagc	accaaactct	tccatgttnt	tccagagctc	420
ctccagccaa	cccaaatagc	cgctgctatn	gtgtagaaca	tcctctgn		467

&lt;210&gt; 76

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(400)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 76

aagctgacag	cattcggggc	gagatgtctc	gctccgtggc	cttagctgtg	ctcgcgctac	60
tctctctttc	tggcctggag	gctatccagc	gtactccaaa	gattcagggt	tactcacgtc	120
atccagcaga	gaatggaaa	tcaaatttcc	tgaattgcta	tgtgtctggg	tttcatccat	180
ccgacattga	agttgactta	ctgaagaatg	gagagagaat	tgaaaaagtg	gagcattcag	240
acttgctttt	cagcaaggac	tggctcttct	atctcttgta	ctacactgaa	ttcaccctca	300
ctgaaaaaga	tgagtatgcc	tgccgtgtga	accatgtgac	tttgtcacag	cccaagatng	360
ttnagtggga	tcganacatg	taagcagcan	catggggaggt			400

&lt;210&gt; 77

&lt;211&gt; 248

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 77

ctggagtgcc	ttgggtgttc	aagccccctgc	aggaagcaga	atgcaccttc	tgaggcacct	60
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ccagctgccc	cggcggggga	tgcgaggctc	ggagcaccct	tgcccggctg	tgattgctgc	120
caggcactgt	tcattctcagc	ttttctgtcc	ctttgctccc	ggcaagcgct	tctgctgaaa	180
gttcatatct	ggagcctgat	gtcttaacga	ataaaggctc	catgctccac	ccgaaaaaaa	240
aaaaaaaa						248

&lt;210&gt; 78

&lt;211&gt; 201

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 78

actagtccag	tgtggtggaa	ttccattgtg	ttgggcccac	cacaatggct	acctttaaca	60
tcacccagac	cccgccctgc	ccgtgcccc	cgctgctgct	aacgacagta	tgatgcttac	120
tctgtacttc	ggaaactatt	tttatgtaat	taatgtatgc	tttcttggtt	ataaatgcct	180
gatttaaaaa	aaaaaaaaaa	a				201

&lt;210&gt; 79

&lt;211&gt; 552

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(552)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 79

tccttttgtt	aggtttttga	gacaacccta	gacctaaact	gtgtcacaga	cttctgaatg	60
tttaggcagt	gctagtaatt	tcctcgtaat	gattctgtta	ttactttcct	attctttatt	120
cctctttctt	ctgaagatta	atgaagtga	aaattgaggt	ggataaatac	aaaaaggtag	180
tgtgatagta	taagtatcta	agtgcagatg	aaagtgtgtt	atatatatcc	attcaaaatt	240
atgcaagtta	gtaattactc	agggttaact	aaattacttt	aatatgctgt	tgaacctact	300
ctgttccttg	gctagaaaaa	attataaaca	ggactttgtt	agtttgggaa	gccaaattga	360
taatattcta	tgttctaaaa	gttgggctat	acataaanta	tnaagaaata	tggaatttta	420
ttcccaggaa	tatgggggtt	atztatgaat	antacccggg	anagaagttt	tgantnaaac	480
cngttttggt	taatacggtta	atatgtcctn	aatnaacaag	gcntgactta	tttccaaaaa	540
aaaaaaaaaa	aa					552

&lt;210&gt; 80

&lt;211&gt; 476

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(476)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 80

acagggattt	gagatgctaa	ggccccagag	atcgtttgat	ccaaccctct	tattttcaga	60
ggggaaaaatg	gggcctagaa	gttacagagc	atctagctgg	tgcgctggca	cccctggcct	120
cacacagact	cccagagtagc	tgggactaca	ggcacacagt	cactgaagca	ggccctgttt	180
gcaattcacg	ttgccacctc	caacttaaac	attcttcata	tgtgatgtcc	ttagtcacta	240
aggttaaact	ttcccaccca	gaaaaggcaa	cttagataaa	atcttagagt	actttcatac	300
tcttctaagt	cctcttccag	cctcactttg	agtcctcctt	gggggttgat	aggaantntc	360

tcttggtttt ctcaataaaa tctctatcca tctcatgttt aatttggtac gcntaaaaat 420  
gctgaaaaaa ttaaaatggt ctggtttcnc tttaaaaaaa aaaaaaaaaa aaaaaa 476

<210> 81  
<211> 232  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(232)  
<223> n = A,T,C or G

<400> 81  
tttttttttg tatgcctnctn ctgtggngtt attgttgctg ccaccctgga ggagcccagt 60  
ttcttctgta tctttctttt ctggggggtc ttcttggtc tgccctcca ttcccagcct 120  
ctcatcccca tcttgcaatt ttgctagggt tggagcgct ttctggtag cccctcagag 180  
actcagtcag cgggaataag tcttaggggt ggggggtgtg gcaagccggc ct 232

<210> 82  
<211> 383  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(383)  
<223> n = A,T,C or G

<400> 82  
aggcgggagc agaagctaaa gccaaagccc aagaagagtg gcagtgccag cactggtgcc 60  
agtaccagta ccaataacat gccagtgcc gtgccagcac cagtgggtggc ttcagtgtgt 120  
gtgccagcct gaccgccact ctcacatttg ggctcttcgc tggccttggt ggagctggtg 180  
ccagcaccag tggcagctct ggtgcctgtg gtttctccta caagtgagat tttagatatt 240  
gttaatcctg ccagtctttc tcttcaagcc aggggtgcac ctcagaaacc tactcaacac 300  
agcactctng gcagccacta tcaatcaatt gaagttgaca ctctgcatta aatctatttg 360  
ccatttcaaa aaaaaaaaaa aaa 383

<210> 83  
<211> 494  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(494)  
<223> n = A,T,C or G

<400> 83  
accgaattgg gaccgtggc ttataagcga tcatgtctc cagtattacc tcaacgagca 60  
gggagatcga gtctatacgc tgaagaaatt tgaccgatg ggacaacaga cctgctcagc 120  
ccatcctgct cgttctccc agatgacaa atactctcga caccgaatca ccatcaagaa 180  
acgcttcaag gtgctcatga cccagcaacc gcgcctgtc ctctgagggt ccttaaaactg 240  
atgtcttttc tgccacctgt taccctcgg agactccgta accaaactct tcggactgtg 300  
agccctgatg cctttttgcc agccatactc tttggentcc agtctctcgt ggcgattgat 360

tatgcttgtg	tgaggcaatc	atggtggcat	cacccatnaa	gggaacacat	ttganttttt	420
tttcncatat	tttaaattac	naccagaata	nttcagaata	aatgaattga	aaaactctta	480
aaaaaaaaaa	aaaa					494

<210> 84  
 <211> 380  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(380)  
 <223> n = A,T,C or G

<400> 84	
gctggtagcc	60
atggtggcat	120
cacccatnaa	180
gggaacacat	240
ttganttttt	300
tttcncatat	360
tttaaattac	380

<210> 85  
 <211> 481  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(481)  
 <223> n = A,T,C or G

<400> 85	
gagttagctc	60
tnccatcgtc	120
ggaaactctc	180
tgtgaaagga	240
gtcgattctg	300
ctatcatgcc	360
ccagattctg	420
aaagaacacc	480
t	481

<210> 86  
 <211> 472  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(472)  
 <223> n = A,T,C or G

<400> 86

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aacatcttcc tgtataatgc tgtgtaatat cgatccgatn ttgtctgctg agaattcatt      60
acttgaaaaa gcaacttnaa gcctggacac tggattataa attcacaata tgcaacactt      120
taaacagtgt gtcaatctgc tcccttactt tgtcatcacc agtctgggaa taagggtatg      180
ccctattcac acctgttaaa agggcgctaa gcatttttga ttcaacatct ttttttttga      240
cacaagtccg aaaaaagcaa aagtaaacag ttnttaattt gttagccaat tcactttctt      300
catgggacag agccatttga tttaaaaagc aaattgcata atattgagct ttgggagctg      360
atatntgagc ggaagantag cctttctact tcaccagaca caactccttt catattggga      420
tgttnacnaa agttatgtct cttacagatg ggatgctttt gtggcaattc tg              472

```

```

<210> 87
<211> 413
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(413)
<223> n = A,T,C or G

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```

<400> 87
agaaaccagt atctctnaaa acaacctctc ataccttgtg gacctaattt tgtgtgctg      60
tgtgtgtgctg cgcatattat atagacaggc acatcttttt tacttttgta aaagcttatg      120
cctcttttgg atctatatct gtgaaagttt taatgatctg ccataatgtc ttggggacct      180
ttgtcttctg tgtaaatggt actagagaaa acacctatnt tatgagtcaa tctagttngt      240
tttattcgac atgaaggaaa tttccagatn acaacactna caaactctcc cttgactagg      300
ggggacaaaag aaaagcnaaa ctgaacatna gaaacaattn cctggtgaga aattncataa      360
acagaaattg ggtingtatat tgaaananng catcattnaa acgttttttt ttt              413

```

```

<210> 88
<211> 448
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(448)
<223> n = A,T,C or G

```

```

<400> 88
cgcagcgggt cctctctatc tagctccagc ctctcgctg cccactccc cgcgtcccgc      60
gtcctagccn accatggccg ggcccctgcg cgcccgcgtg ctctgctgg ccactctggc      120
cgtggccctg gccgtgagcc cgcggccggt ctccagtcgc ggcaagccgc cgcgcctggg      180
gggaggccca tggaccccgc gtggaagaag aagggtgtgc gcgtgcactg gactttgccg      240
tcggcnanta caacaaaccc gcaacnactt ttaccnagcn cgcgtgcag gttgtgccgc      300
cccaancaaa ttgttactng gggtaantaa ttcttggaag ttgaacctgg gccaaaacng      360
tttaccagaa ccnagccaat tngaacaatt nccctccat aacagcccct tttaaaaagg      420
gaancantcc tgntcttttc caaatttt              448

```

```

<210> 89
<211> 463
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature

```

&lt;222&gt; (1)...(463)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 89

gaattttgtg	cactggccac	tgtgatggaa	ccattgggcc	aggatgcttt	gagtttatca	60
gtagtgattc	tgccaaagtt	ggtgttgtaa	catgagtatg	taaaatgtca	aaaaattagc	120
agaggtctag	gtctgcatat	cagcagacag	tttgcccggtg	tattttgtag	ccttgaagtt	180
ctcagtgaca	agttntttct	gatgcgaagt	tctnattcca	gtgttttagt	cctttgcac	240
tttnatgtn	agacttgctt	ctntnaaatt	gcttttgtnt	tctgcaggta	ctatctgtgg	300
tttaacaaaa	tagaannact	tctctgcttn	gaanatttga	atatcttaca	tctnaaaatn	360
aattctctcc	ccatannaaa	acccangccc	ttggganaat	ttgaaaaang	gntccttcnn	420
aattcnnana	anttcagntn	tcatacaaca	naacngganc	ccc		463

&lt;210&gt; 90

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(400)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 90

agggattgaa	ggtctnttnt	actgtcggac	tgttcancca	ccaactctac	aagttgctgt	60
cttccactca	ctgtctgtaa	gcntnttaac	ccagactgta	tcttcataaa	tagaacaaat	120
tcttcaccag	tcacatcttc	taggaccttt	ttggattcag	ttagtataag	ctcttccact	180
tcctttgtta	agacttcate	tggtaaagtc	ttaagttttg	tagaaaggaa	tttaattgct	240
cgttctctaa	caatgtcctc	tccttgaagt	atttggctga	acaaccacc	tnaagtcctt	300
ttgtgcatcc	attttaaata	tacttaatag	ggcattggtn	cactagggtta	aattctgcaa	360
gagtcactctg	tctgcaaaag	ttgcgttagt	atatctygca			400

&lt;210&gt; 91

&lt;211&gt; 480

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(480)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 91

gagctcggat	ccaataatct	ttgtctgagg	gcagcacaca	tatncagtgc	catggnaact	60
ggtctacccc	acatgggagc	agcatgccgt	agntatataa	ggtcattccc	tgagtcagac	120
atgcctcttt	gactaccgtg	tgccagtgtc	ggtgattctc	acacacctcc	nnccgtcttt	180
tgtggaaaaa	ctggcacttg	nctggaacta	gcaagacatc	acttacaaat	tcaccacaga	240
gacacttgaa	aggtgtaaca	aagcgactct	tgcattgctt	tttgtccctc	cggcaccagt	300
tgtcaatact	aacccgctgg	tttgccctcca	tcacatttgt	gatctgtagc	tctggatata	360
tctcctgaca	gtactgaaga	acttcttctt	ttgtttcaaa	agcaactctt	ggtgcctgtt	420
ngatcagggt	cccatttccc	agtccgaatg	ttcacatggc	atatnttact	tcccacaaaa	480

&lt;210&gt; 92

&lt;211&gt; 477

&lt;212&gt; DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(477)

<223> n = A,T,C or G

<400> 92

atacagccca	natcccacca	cgaagatgcg	cttggttgact	gagaacctga	tgcgggtcact	60
ggccccgctg	tagccccagc	gactctccac	ctgctggaag	cggttgatgc	tgcactcctt	120
cccacgcagg	cagcagcggg	gccgggtcaat	gaactccact	cgtggccttg	ggttgacggg	180
taantgcagg	aagaggctga	ccacctcgcg	gtccaccagg	atgcccgact	gtgcggggacc	240
tgcagcgaaa	ctcctcgatg	gtcatgagcg	ggaagcgaat	gangcccagg	gccttgccca	300
gaaccttccg	cctgtttctt	ggcgtcacct	gcagctgctg	ccgctnacac	tgggcctcgg	360
accagcggac	aaacggcggt	gaacagccgc	acctcacgga	tgcccantgt	gtcgcgctcc	420
aggaacggcn	ccagcgtgtc	caggtcaatg	tcggtgaanc	ctccgcgggt	aatggcg	477

<210> 93

<211> 377

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(377)

<223> n = A,T,C or G

<400> 93

gaacggctgg	accttgccctc	gcattgtgct	gctggcagga	ataccttggc	aagcagctcc	60
agtccgagca	gccccagacc	gctgccgccc	gaagctaagc	ctgcctcttg	ccttcccctc	120
cgcctcaatg	cagaaccant	agtgggagca	ctgtgttttag	agttaagagt	gaacactgtg	180
tgattttact	tgggaatttc	ctctgttata	tagcttttcc	caatgctaata	ttccaaacaa	240
caacaacaaa	ataacatgtt	tgctgtttna	gttggtataaa	agtangtgat	tctgtatnta	300
aagaaaatat	tactgttaca	tatactgctt	gcaanttctg	tatttattgg	tnctctggaa	360
ataaatatat	tattaaa					377

<210> 94

<211> 495

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(495)

<223> n = A,T,C or G

<400> 94

ccctttgagg	ggttaggggc	cagttcccag	tggaagaaac	agggcaggag	aantgcgtgc	60
cgagctgang	cagatttccc	acagtgaccc	cagagccctg	ggctatagtc	tctgaccctt	120
ccaaggaaag	accaccttct	ggggacatgg	gctggagggc	aggacctaga	ggcaccaagg	180
gaaggcccca	ttccggggct	gttccccgag	gaggaagggg	aggggctctg	tgtgcccccc	240
acgaggaana	ggccctgant	cctgggatca	nacaccctt	cacgtgtatc	cccacacaaa	300
tgcaagctca	ccaaggtccc	ctctcagtc	cttccctaca	ccctgaacgg	ncactggccc	360
acacccaccc	agancancca	cccgccatgg	ggaatgtnt	caaggaatcg	cngggcaacg	420
tggactctng	tcccnnaagg	gggcagaatc	tccaatagan	gganngaacc	cttgctnana	480

aaaaaaaaana aaaaaa

495

<210> 95  
 <211> 472  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(472)  
 <223> n = A,T,C or G

&lt;400&gt; 95

ggttacttgg	tttcattgcc	accacttagt	ggatgtcatt	tagaaccatt	ttgtctgctc	60
cctctggaag	ccttgccgag	agcggacttt	gtaattgttg	gagaataact	gctgaatttt	120
tagctgtttt	gagttgattc	gcaccactgc	accacaactc	aatatgaaaa	ctatttnact	180
tatttattat	cttgtgaaaa	gtatacaatg	aaaattttgt	tcatactgta	tttatcaagt	240
atgatgaaaa	gcaatagata	tatattcttt	tattatgttn	aattatgatt	gccattatta	300
atcggcaaaa	tgtggagtgt	atgttctttt	cacagtaata	tatgcctttt	gtaacttcac	360
ttggttattt	tattgtaa	gaattacaaa	attcttaatt	taagaaaatg	gtangttata	420
tttanttcan	taatttcttt	ccttggtttac	gttaattttg	aaaagaatgc	at	472

<210> 96  
 <211> 476  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(476)  
 <223> n = A,T,C or G

&lt;400&gt; 96

ctgaagcatt	tcttcaaact	tntctacttt	tgtcattgat	acctgtagta	agttgacaat	60
gtggtgaaat	ttcaaaatta	tatgtaactt	ctactagttt	tactttctcc	cccaagtctt	120
ttttaactca	tgattttttac	acacacaatc	cagaacttat	tatatagcct	ctaagtcttt	180
attcttcaca	gtagatgatg	aaagagtctt	ccagtgtctt	gngcanaatg	ttctagntat	240
agctggatac	atacngtggg	agttctataa	actcatacct	cagtgggact	naaccaaaat	300
tgtgttagtc	tcaattccta	ccacactgag	ggagcctccc	aatcactat	attcttatct	360
gcaggtactc	ctccagaaaa	acngacaggg	caggcttgca	tgaaaaagtn	acatctgcgt	420
tacaaagtct	atcttcctca	nangtctgtn	aaggaacaat	ttaatcttct	agcttt	476

<210> 97  
 <211> 479  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(479)  
 <223> n = A,T,C or G

&lt;400&gt; 97

actctttcta	atgctgatat	gatcttgagt	ataagaatgc	atatgtcact	agaatggata	60
aaataatgct	gcaaacttaa	tgttcttatg	caaaatggaa	cgctaataa	acacagctta	120



caatcgcaaa tcaaaactca caagtgtca tctgtttag atttagtgta ataagactta	180
gattgtgtc cttcggatat gattgtttct canatcttgg gcaatnttcc ttagtcaaat	240
caggctacta gaattctgtt attggatatn tgagagcatg aaatttttaa naatacactt	300
gtgattatna aattaatcac aaatttcact tatacctgct atcagcagct agaaaaacat	360
ntnnttttta natcaaagta ttttgtgttt ggaantgtnn aaatgaaatc tgaatgtggg	420
ttcnatctta ttttttcccn gacnactant tnttttttta gggncatttc tganccatc	479

&lt;210&gt; 98

&lt;211&gt; 461

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 98

agtgacttgt cctccaacaa aacccttga tcaagtttgt ggcactgaca atcagacctta	60
tgctagtcc tgcctctat tcgtactaa atgcagactg gaggggacca aaaaggggca	120
tcaactccag ctggattatt ttggagcctg caaatctatt cctacttgta cggactttga	180
agtgattcag tttcctctac ggatgagaga ctggctcaag aatctctca tgcagcttta	240
tgaagccact ctgaacacgc tggttatcta gatgagaaca gagaaataaa gtcagaaaaat	300
ttacctggag aaaagaggct ttggctgggg accatcccat tgaaccttct cttaaggact	360
ttaagaaaaa ctaccacatg ttgtgtatcc tggtgccggc cgtttatgaa ctgaccaccc	420
tttggaataa tcttgacgct cctgaacttg ctctctgcg a	461

&lt;210&gt; 99

&lt;211&gt; 171

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 99

gtggccgcgc gcaggtgttt cctcgtaccg cagggccccc tcccttcccc aggcgtccct	60
cggcgctct gcgggcccga ggaggagcgg ctggcggtg gggggagtgt gaccaccct	120
cggtgagaaa agccttctct agcgatctga gaggcgtgcc ttgggggtac c	171

&lt;210&gt; 100

&lt;211&gt; 269

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 100

cggccgcaag tgcaactcca gctggggccg tgcggacgaa gattctgcc a gcagttggtc	60
cgactgcgac gacggcggcg gcgacagtcg caggtgcagc gcgggcgcct ggggtcttg	120
aaggctgagc tgacgccgca gaggtcgtgt caggtccac gaccttgacg ccgtcgggga	180
cagccggaac agagcccggg gaagcgggag gcctcgggga gcccctcggg aagggcggcc	240
cgagagatac gcaggtgcag gtggccgcc	269

&lt;210&gt; 101

&lt;211&gt; 405

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 101

tttttttttt ttttgaatc tactgcgagc acagcaggtc agcaacaagt ttattttgca	60
gctagcaagg taacagggtg gggcatgggt acatgttcag gtcaacttcc tttgtcgtgg	120
ttgattgggt tgtctttatg ggggcggggg ggggtagggg aaacgaagca aataacatgg	180
agtgggtgca cctccctgt agaacctgt tacaaagctt ggggcagttc acctggtctg	240
tgacctcat tttcttgaca tcaatgttat tagaagtcag gatattttt agagagtcca	300

ctgttctgga gggagattag ggtttcttgc caaatccaac aaaatccact gaaaaagttg 360  
 gatgatcagt acgaataccg aggcattatc tcatatcggt ggcca 405

<210> 102  
 <211> 470  
 <212> DNA  
 <213> Homo sapien

<400> 102  
 tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt 60  
 ggcacttaat ccatttttat ttcaaaatgt ctacaaattt aatcccatta tacggtattt 120  
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 <213> Homo sapien

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 <212> DNA  
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 <211> 538  
 <212> DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 105

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&lt;210&gt; 106

&lt;211&gt; 473

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 106

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&lt;210&gt; 107

&lt;211&gt; 1621

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 107

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a 1621

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<210> 108  
 <211> 382  
 <212> PRT  
 <213> Homo sapien

<400> 108

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35     40     45
Gly Lys Arg Ser Leu Val Leu Asp Leu Lys Gln Pro Arg Gly Ala Ala
50     55     60
Val Leu Arg Arg Leu Cys Lys Arg Ser Asp Val Leu Leu Glu Pro Phe
65     70     75     80
Arg Arg Gly Val Met Glu Lys Leu Gln Leu Gly Pro Glu Ile Leu Gln
85     90     95
Arg Glu Asn Pro Arg Leu Ile Tyr Ala Arg Leu Ser Gly Phe Gly Gln
100    105    110
Ser Gly Ser Phe Cys Arg Leu Ala Gly His Asp Ile Asn Tyr Leu Ala
115    120    125
Leu Ser Gly Val Leu Ser Lys Ile Gly Arg Ser Gly Glu Asn Pro Tyr
130    135    140
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145    150    155    160
Ala Leu Gly Ile Ile Met Ala Leu Phe Asp Arg Thr Arg Thr Asp Lys
165    170    175
Gly Gln Val Ile Asp Ala Asn Met Val Glu Gly Thr Ala Tyr Leu Ser
180    185    190
Ser Phe Leu Trp Lys Thr Gln Lys Ser Ser Leu Trp Glu Ala Pro Arg
195    200    205
Gly Gln Asn Met Leu Asp Gly Gly Ala Pro Phe Tyr Thr Thr Tyr Arg
210    215    220
Thr Ala Asp Gly Glu Phe Met Ala Val Gly Ala Ile Glu Pro Gln Phe
225    230    235    240
Tyr Glu Leu Leu Ile Lys Gly Leu Gly Leu Lys Ser Asp Glu Leu Pro
245    250    255
Asn Gln Met Ser Met Asp Asp Trp Pro Glu Met Lys Lys Lys Phe Ala
260    265    270
Asp Val Phe Ala Lys Lys Thr Lys Ala Glu Trp Cys Gln Ile Phe Asp
275    280    285
Gly Thr Asp Ala Cys Val Thr Pro Val Leu Thr Phe Glu Glu Val Val
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His His Asp His Asn Lys Glu Arg Gly Ser Phe Ile Thr Ser Glu Glu
305    310    315    320
Gln Asp Val Ser Pro Arg Pro Ala Pro Leu Leu Leu Asn Thr Pro Ala

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325 330 335  
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 <211> 1524  
 <212> DNA  
 <213> Homo sapien

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 <211> 3410  
 <212> DNA  
 <213> Homo sapien

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&lt;210&gt; 111

&lt;211&gt; 1289

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 111

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&lt;210&gt; 112

&lt;211&gt; 315

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 112

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Met Val Phe Thr Val Arg Leu Leu His Ile Phe Thr Val Asn Lys Gln
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Leu Gly Pro Lys Ile Val Ile Val Ser Lys Met Met Lys Asp Val Phe
          20          25          30
Phe Phe Leu Phe Phe Leu Gly Val Trp Leu Val Ala Tyr Gly Val Ala
          35          40          45
Thr Glu Gly Leu Leu Arg Pro Arg Asp Ser Asp Phe Pro Ser Ile Leu
          50          55          60
Arg Arg Val Phe Tyr Arg Pro Tyr Leu Gln Ile Phe Gly Gln Ile Pro
          65          70          75          80
Gln Glu Asp Met Asp Val Ala Leu Met Glu His Ser Asn Cys Ser Ser
          85          90          95
Glu Pro Gly Phe Trp Ala His Pro Pro Gly Ala Gln Ala Gly Thr Cys
          100          105          110
Val Ser Gln Tyr Ala Asn Trp Leu Val Val Leu Leu Leu Val Ile Phe
          115          120          125
Leu Leu Val Ala Asn Ile Leu Leu Val Asn Leu Leu Ile Ala Met Phe
          130          135          140
Ser Tyr Thr Phe Gly Lys Val Gln Gly Asn Ser Asp Leu Tyr Trp Lys
          145          150          155          160
Ala Gln Arg Tyr Arg Leu Ile Arg Glu Phe His Ser Arg Pro Ala Leu
          165          170          175
Ala Pro Pro Phe Ile Val Ile Ser His Leu Arg Leu Leu Leu Arg Gln
          180          185          190
Leu Cys Arg Arg Pro Arg Ser Pro Gln Pro Ser Ser Pro Ala Leu Glu

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195                      200                      205  
 His Phe Arg Val Tyr Leu Ser Lys Glu Ala Glu Arg Lys Leu Leu Thr  
 210                      215                      220  
 Trp Glu Ser Val His Lys Glu Asn Phe Leu Leu Ala Arg Ala Arg Asp  
 225                      230                      235                      240  
 Lys Arg Glu Ser Asp Ser Glu Arg Leu Lys Arg Thr Ser Gln Lys Val  
 245                      250                      255  
 Asp Leu Ala Leu Lys Gln Leu Gly His Ile Arg Glu Tyr Glu Gln Arg  
 260                      265                      270  
 Leu Lys Val Leu Glu Arg Glu Val Gln Gln Cys Ser Arg Val Leu Gly  
 275                      280                      285  
 Trp Val Ala Glu Ala Leu Ser Arg Ser Ala Leu Leu Pro Pro Gly Gly  
 290                      295                      300  
 Pro Pro Pro Pro Asp Leu Pro Gly Ser Lys Asp  
 305                      310                      315

&lt;210&gt; 113

&lt;211&gt; 553

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 113

Met Val Gln Arg Leu Trp Val Ser Arg Leu Leu Arg His Arg Lys Ala  
 1                      5                      10                      15  
 Gln Leu Leu Leu Val Asn Leu Leu Thr Phe Gly Leu Glu Val Cys Leu  
 20                      25                      30  
 Ala Ala Gly Ile Thr Tyr Val Pro Pro Leu Leu Leu Glu Val Gly Val  
 35                      40                      45  
 Glu Glu Lys Phe Met Thr Met Val Leu Gly Ile Gly Pro Val Leu Gly  
 50                      55                      60  
 Leu Val Cys Val Pro Leu Leu Gly Ser Ala Ser Asp His Trp Arg Gly  
 65                      70                      75                      80  
 Arg Tyr Gly Arg Arg Arg Pro Phe Ile Trp Ala Leu Ser Leu Gly Ile  
 85                      90                      95  
 Leu Leu Ser Leu Phe Leu Ile Pro Arg Ala Gly Trp Leu Ala Gly Leu  
 100                      105                      110  
 Leu Cys Pro Asp Pro Arg Pro Leu Glu Leu Ala Leu Leu Ile Leu Gly  
 115                      120                      125  
 Val Gly Leu Leu Asp Phe Cys Gly Gln Val Cys Phe Thr Pro Leu Glu  
 130                      135                      140  
 Ala Leu Leu Ser Asp Leu Phe Arg Asp Pro Asp His Cys Arg Gln Ala  
 145                      150                      155                      160  
 Tyr Ser Val Tyr Ala Phe Met Ile Ser Leu Gly Gly Cys Leu Gly Tyr  
 165                      170                      175  
 Leu Leu Pro Ala Ile Asp Trp Asp Thr Ser Ala Leu Ala Pro Tyr Leu  
 180                      185                      190  
 Gly Thr Gln Glu Glu Cys Leu Phe Gly Leu Leu Thr Leu Ile Phe Leu  
 195                      200                      205  
 Thr Cys Val Ala Ala Thr Leu Leu Val Ala Glu Glu Ala Ala Leu Gly  
 210                      215                      220  
 Pro Thr Glu Pro Ala Glu Gly Leu Ser Ala Pro Ser Leu Ser Pro His  
 225                      230                      235                      240  
 Cys Cys Pro Cys Arg Ala Arg Leu Ala Phe Arg Asn Leu Gly Ala Leu  
 245                      250                      255  
 Leu Pro Arg Leu His Gln Leu Cys Cys Arg Met Pro Arg Thr Leu Arg



260 265 270  
 Arg Leu Phe Val Ala Glu Leu Cys Ser Trp Met Ala Leu Met Thr Phe  
 275 280 285  
 Thr Leu Phe Tyr Thr Asp Phe Val Gly Glu Gly Leu Tyr Gln Gly Val  
 290 295 300  
 Pro Arg Ala Glu Pro Gly Thr Glu Ala Arg Arg His Tyr Asp Glu Gly  
 305 310 315 320  
 Val Arg Met Gly Ser Leu Gly Leu Phe Leu Gln Cys Ala Ile Ser Leu  
 325 330 335  
 Val Phe Ser Leu Val Met Asp Arg Leu Val Gln Arg Phe Gly Thr Arg  
 340 345 350  
 Ala Val Tyr Leu Ala Ser Val Ala Ala Phe Pro Val Ala Ala Gly Ala  
 355 360 365  
 Thr Cys Leu Ser His Ser Val Ala Val Val Thr Ala Ser Ala Ala Leu  
 370 375 380  
 Thr Gly Phe Thr Phe Ser Ala Leu Gln Ile Leu Pro Tyr Thr Leu Ala  
 385 390 395 400  
 Ser Leu Tyr His Arg Glu Lys Gln Val Phe Leu Pro Lys Tyr Arg Gly  
 405 410 415  
 Asp Thr Gly Gly Ala Ser Ser Glu Asp Ser Leu Met Thr Ser Phe Leu  
 420 425 430  
 Pro Gly Pro Lys Pro Gly Ala Pro Phe Pro Asn Gly His Val Gly Ala  
 435 440 445  
 Gly Gly Ser Gly Leu Leu Pro Pro Pro Pro Ala Leu Cys Gly Ala Ser  
 450 455 460  
 Ala Cys Asp Val Ser Val Arg Val Val Val Gly Glu Pro Thr Glu Ala  
 465 470 475 480  
 Arg Val Val Pro Gly Arg Gly Ile Cys Leu Asp Leu Ala Ile Leu Asp  
 485 490 495  
 Ser Ala Phe Leu Leu Ser Gln Val Ala Pro Ser Leu Phe Met Gly Ser  
 500 505 510  
 Ile Val Gln Leu Ser Gln Ser Val Thr Ala Tyr Met Val Ser Ala Ala  
 515 520 525  
 Gly Leu Gly Leu Val Ala Ile Tyr Phe Ala Thr Gln Val Val Phe Asp  
 530 535 540  
 Lys Ser Asp Leu Ala Lys Tyr Ser Ala  
 545 550

&lt;210&gt; 114

&lt;211&gt; 241

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 114

Met Gln Cys Phe Ser Phe Ile Lys Thr Met Met Ile Leu Phe Asn Leu  
 1 5 10 15  
 Leu Ile Phe Leu Cys Gly Ala Ala Leu Leu Ala Val Gly Ile Trp Val  
 20 25 30  
 Ser Ile Asp Gly Ala Ser Phe Leu Lys Ile Phe Gly Pro Leu Ser Ser  
 35 40 45  
 Ser Ala Met Gln Phe Val Asn Val Gly Tyr Phe Leu Ile Ala Ala Gly  
 50 55 60  
 Val Val Val Phe Ala Leu Gly Phe Leu Gly Cys Tyr Gly Ala Lys Thr  
 65 70 75 80  
 Glu Ser Lys Cys Ala Leu Val Thr Phe Phe Phe Ile Leu Leu Leu Ile

```
<210> 115
<211> 366
<212> DNA
<213> Homo sapien
```

```
<210> 116
<211> 282
<212> DNA
<213> Homo sapien
```

<400> 116						
acaaagatga	accatttct	atattatagc	aaaattaaaa	tctaccgta	ttctaattatt	60
gagaaatgag	atnaaacaca	atnttataaa	gtctacttag	agaagatcaa	gtgacctcaa	120
agactttact	attttcatat	tttaagacac	atgatttatac	ctatttttagt	aacctgggtc	180
atacgttaaa	caaaggataa	tgtgaacagc	agagaggatt	tggtggcaga	aaatctatgt	240
tcaatctnga	actatctana	tcacagacat	ttctatttct	tt		282

<210> 117  
<211> 305

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(305)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 117

acacatgtcg	cttcactgcc	ttcttagatg	cttctgggtca	acatanagga	acagggacca	60
tatttatcct	ccctcctgaa	acaattgcaa	aataanacaa	aatatatgaa	acaattgcaa	120
aataaggcaa	aatatatgaa	acaacagggtc	tcgagatatt	ggaaatcagt	caatgaagga	180
tactgatccc	tgatcactgt	cctaattgcag	gatgtgggaa	acagatgagg	tcacctctgt	240
gactgccccca	gcttactgcc	tgtagagagt	ttctangctg	cagttcagac	agggagaaat	300
tggggt						305

&lt;210&gt; 118

&lt;211&gt; 71

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(71)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 118

accaaggtgt	ntgaatctct	gacgtgggga	tctctgattc	cgcacaatc	tgagtggaaa	60
aantcctggg	t					71

&lt;210&gt; 119

&lt;211&gt; 212

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(212)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 119

actccggttg	gtgtcagcag	cacgtggcat	tgaacatngc	aatgtggagc	ccaaaccaca	60
gaaaatgggg	tgaaattggc	caactttcta	tnaacttatg	ttggcaantt	tgccaccaac	120
agtaagctgg	cccttcta	aaaagaaaat	tgaaaggttt	ctcactaanc	ggaattaant	180
aatggantca	aganactccc	aggcctcagc	gt			212

&lt;210&gt; 120

&lt;211&gt; 90

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(90)

&lt;223&gt; n = A,T,C or G

<400> 120  
 actcgttgca natcaggggc cccccagagt caccgttgca ggagtccttc tggctcttgcc 60  
 ctccgccggc gcagaacatg ctgggggtggt 90

<210> 121  
 <211> 218  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(218)  
 <223> n = A,T,C or G

<400> 121  
 tgtancgtga anacgacaga nagggttgct aaaaatggag aanccttgaa gtcattttga 60  
 gaataagatt tgctaaaaga tttggggcta aaacatgggtt attgggagac atttctgaag 120  
 atatncangt aaattangga atgaattcat ggttcttttg ggaattcctt tacgatngcc 180  
 agcatanact tcatgtgggg atancagcta cccttgta 218

<210> 122  
 <211> 171  
 <212> DNA  
 <213> Homo sapien

<400> 122  
 taggggtgta tgcaactgta aggacaaaaa ttgagactca actggcttaa ccaataaagg 60  
 catttgtagt ctcatggaac aggaagtcgg atgggtggggc atcttcagtg ctgcatgagt 120  
 caccaccccg gcgggggtcat ctgtgccaca ggtccctggt gacagtgcgg t 171

<210> 123  
 <211> 76  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(76)  
 <223> n = A,T,C or G

<400> 123  
 tgtagcgtga agacnacaga atgggtgtgtg ctgtgctatc caggaacaca tttattatca 60  
 ttatcaanta ttgtgt 76

<210> 124  
 <211> 131  
 <212> DNA  
 <213> Homo sapien

<400> 124  
 acctttcccc aaggccaatg tcctgtgtgc taactggccg gctgcaggac agctgcaatt 60  
 caatgtgctg ggtcatatgg aggggaggag actctaaaat agccaatttt atttctcttg 120  
 ttaagatttg t 131

<210> 125  
 <211> 432  
 <212> DNA  
 <213> Homo sapien

<400> 125  
 accttatcta ctggctatga aatagatggt ggaaaattgc gttaccaact ataccactgg 60  
 cttgaaaaag aggtgatagc tcttcagagg acttgtagt tttgctcaga tgctgaagaa 120  
 ctacagtctg catttggcag aaatgaagat gaatttggat taaatgagga tgctgaagat 180  
 ttgcctcacc aaacaaaagt gaaacaactg agagaaaatt ttcaggaaaa aagacagtgg 240  
 ctcttgaagt atcagtcact tttgagaatg tttcttagtt actgcatact tcatggatcc 300  
 catgggtgggg gtcttgcacg tgtaagaatg gaattgattt tgcttttgca agaattctcag 360  
 caggaaacat cagaaccact attttctagc cctctgtcag agcaaacctc agtgcctctc 420  
 ctctttgctt gt 432

<210> 126  
 <211> 112  
 <212> DNA  
 <213> Homo sapien

<400> 126  
 acacaacttg aatagtaaaa tagaaactga gctgaaattt ctaattcact ttctaaccat 60  
 agtaagaatg atatttcccc ccagggatca ccaaatattt ataaaaattt gt 112

<210> 127  
 <211> 54  
 <212> DNA  
 <213> Homo sapien

<400> 127  
 accacgaaac cacaacaag atggaagcat caatccactt gccaaagcaca gcag 54

<210> 128  
 <211> 323  
 <212> DNA  
 <213> Homo sapien

<400> 128  
 acctcattag taattgtttt gttgtttcat ttttttctaa tgtctcccct ctaccagctc 60  
 acctgagata acagaatgaa aatggaagga cagccagatt tctcctttgc tctctgctca 120  
 ttctctctga agtctagggt acccattttg gggaccatt ataggcaata aacacagtgc 180  
 ccaaagcatt tggacagtgt cttgttgtgt tttagaatgg ttttcctttt tcttagcctt 240  
 ttcttgcaaa aggtcactc agtccttgc ttgctcagtg gactgggctc ccaggggcct 300  
 aggtgcctt cttttccatg tcc 323

<210> 129  
 <211> 192  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)... (192)  
 <223> n = A,T,C or G

&lt;400&gt; 129

```

acatacatgt gtgtatatatt ttaaatatca cttttgtatc actctgactt tttagcatatc      60
tgaaaacaca ctaacataat ttntgtgaac catgatcaga tacaacccaa atcattcatc      120
tagcacattc atctgtgata naaagatagg tgagtttcat ttccttcacg ttggccaatg      180
gataaacaat gt                                     192

```

&lt;210&gt; 130

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(362)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 130

```

ccctttttta tggaatgagt agactgtatg tttgaanatt tanccacaac ctctttgaca      60
tataatgacg caacaaaaag gtgctgttta gtcctatggt tcagtttatg cccctgacaa      120
gtttccattg tgttttgccg atcttctggc taatcgtggg atcctccatg ttattagtaa      180
ttctgtattc cattttgcta acgcctggta gatgtaacct gctangaggc taactttata      240
cttattttaa agctcttatt ttgtggatcat taaaatggca atttatgtgc agcactttat      300
tgcagcagga agcacgtgtg ggttggttgt aaagctcttt gctaatttta aaaagtaatg      360
gg                                     362

```

&lt;210&gt; 131

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(332)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 131

```

ctttttgaaa gatcgtgtcc actcctgtgg acatcttggt ttaatggagt ttcccatgca      60
gtangactgg tatggttgca gctgtccaga taaaaacatt tgaagagctc caaaatgaga      120
gttctcccag gtgcgacctg ctgctccaag tctcagcagc agcctctttt aggaggcatc      180
ttctgaacta gattaaggca gcttgtaaatt ctgatgtgat ttggtttatt atccaactaa      240
cttccatctg ttatcactgg agaaagccca gactcccan gacnggtacg gattgtgggc      300
atanaaggat tgggtgaagc tggcgttgtg gt                                     332

```

&lt;210&gt; 132

&lt;211&gt; 322

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(322)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 132

```

acttttgcca tttgttatat ataaacaatc ttgggacatt ctctgaaaa ctaggtgtcc      60

```

```

agtggctaag agaactcgat ttcaagcaat tctgaaagga aaaccagcat gacacagaat 120
ctcaaattcc caaacagggg ctctgtggga aaaatgaggg aggacctttg tatctcgggt 180
tttagcaagt taaaatgaan atgacaggaa aggcttattt atcaacaaaag agaagagttg 240
ggatgcttct aaaaaaaact ttggtagaga aaataggaat gctnaatcct agggaagcct 300
gtaacaatct acaattggtc ca 322

```

&lt;210&gt; 133

&lt;211&gt; 278

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(278)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 133

```

acaagccttc acaagtttaa ctaaattggg attaatcttt ctgtanttat ctgcataatt 60
cttgtttttc tttccatctg gctcctgggt tgacaatttg tggaaacaac tctattgcta 120
ctatttaaaa aaaatcacia atctttccct ttaagctatg ttnaattcaa actattcctg 180
ctattcctgt tttgtcaaag aaattatatt tttcaaaata tgtntatttg tttgatgggt 240
cccacgaaac actaataaaa accacagaga ccagcctg 278

```

&lt;210&gt; 134

&lt;211&gt; 121

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(121)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 134

```

gtttanaaaa ctgttttagc tccatagagg aaagaatggt aaactttgta ttttaaaaca 60
tgattctctg aggttaaact tggttttcaa atgttatatt tacttgatt ttgcttttgg 120
t 121

```

&lt;210&gt; 135

&lt;211&gt; 350

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(350)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 135

```

acttanaacc atgcctagca catcagaate cctcaaagaa catcagtata atcctatacc 60
atancaagtg gtgactggtt aagcgtgcga caaagggtcag ctggcacatt acttggtgtgc 120
aaacttgata cttttgttct aagtaggaac tagtatacag tncctaggan tggtagtcca 180
gggtgcccc caactcctgc agccgtcct ctgtgccagn ccctgnaagg aactttcgct 240
ccacctcaat caagccctgg gccatgctac ctgcaattgg ctgaacaaac gtttgetgag 300
ttcccaagga tgcaaagcct ggtgctcaac tctgggggcg tcaactcagt 350

```

<210> 136  
 <211> 399  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (399)  
 <223> n = A,T,C or G

<400> 136  
 tgtaccgtga agacgacaga agttgcatgg cagggacagg gcagggccga ggccagggtt 60  
 gctgtgattg tatccgaata ntctctgtga gaaaagataa tgagatgacg tgagcagcct 120  
 gcagacttgt gtctgccttc aanaagccag acaggaaggc cctgcctgcc ttggctctga 180  
 cctggcggcc agccagccag ccacaggtgg gcttcttctt tttgtggtga caacnccaag 240  
 aaaactgcag agggccaggg tcaggtgtna gtgggtangt gaccataaaa caccaggtgc 300  
 tcccaggaac cggggcaaag gccatcccca cctacagcca gcatgccacac tggcgtgatg 360  
 ggtgcagang gatgaagcag ccagntgttc tgctgtggt 399

<210> 137  
 <211> 165  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (165)  
 <223> n = A,T,C or G

<400> 137  
 actggtgtgg tnggggggtga tgctgggtgg anaagttgan gtgacttcan gatggtgtgt 60  
 ggaggaagtg tgtgaacgta gggatgtaga ngttttggcc gtgctaaatg agcttcggga 120  
 ttggctggtc ccactgggtg tcactgtcat tgggtggggt cctgt 165

<210> 138  
 <211> 338  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (338)  
 <223> n = A,T,C or G

<400> 138  
 actcactgga atgccacatt cacaacagaa tcagaggtct gtgaaaacat taatggctcc 60  
 ttaacttctc cagtaagaat cagggacttg aaatggaaac gttaacagcc acatgcccaa 120  
 tgctgggcag tctcccatgc ctccacagt gaaagggctt gagaaaaatc acatccaatg 180  
 tcatgtgttt ccagccacac caaaaggtgc ttggggtgga gggctggggg catananggt 240  
 cangcctcag gaagcctcaa gttccattca gctttgccac tgtacattcc ccatntttaa 300  
 aaaaactgat gccttttttt tttttttttg taaaattc 338

<210> 139  
 <211> 382



&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 139

gggaatcttg	gtttttggca	tctgggttgc	ctatagccga	ggccactttg	acagaacaaa	60
gaaagggact	tcgagtaaga	aggtgattta	cagccagcct	agtgcccga	gtgaaggaga	120
attcaaacag	acctcgatcat	tcctgggtgtg	agcctgggtcg	gtccaccgcc	tatcatctgc	180
atttgcccta	ctcaggtgct	accggactct	ggccccctgat	gtctgtagtt	tcacaggatg	240
ccttattttgt	cttctacacc	ccacagggcc	ccctacttct	tcggatgtgt	ttttaataat	300
gtcagctatg	tgccccatcc	tccttcatgc	cctccctccc	tttctacca	ctgctgagtg	360
gcctggaact	tgtttaaagt	gt				382

&lt;210&gt; 140

&lt;211&gt; 200

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (200)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 140

accaaanctt	ctttctgttg	tgtnngattt	tactataggg	gttnngcttn	ttctaaanat	60
acttttcatt	taacancctt	tgtaagtgt	caggctgcac	tttgctccat	anaattattg	120
ttttcacatt	tcaacttgta	tgtgtttgtc	tcttanagca	ttggtgaaat	cacatatttt	180
atattcagca	taaaggagaa					200

&lt;210&gt; 141

&lt;211&gt; 335

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (335)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 141

actttatttt	caaaacactc	atatgttgca	aaaaacacat	agaaaaataa	agtttggtgg	60
gggtgctgac	taaacttcaa	gtcacagact	tttatgtgac	agattggagc	agggtttggt	120
atgcatgtag	agaacccaaa	ctaatttatt	aaacaggata	gaaacaggct	gtctgggtga	180
aatggttctg	agaaccatcc	aattcacctg	tcagatgctg	atanactagc	tcctcagatg	240
tttttctacc	agttcagaga	tnggttaatg	actanttcca	atggggaaaa	agcaagatgg	300
attcacaac	caagtaattt	taaacaaaga	cactt			335

&lt;210&gt; 142

&lt;211&gt; 459

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (459)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 142

accagggttaa	tattgccaca	tatatccttt	ccaattgcgg	gctaaacaga	cgtgtattta	60
gggttggtta	aagacaaccc	agcttaatat	caagagaaat	tgtgaccttt	catggagtat	120
ctgatggaga	aaacactgag	ttttgacaaa	tcttatttta	ttcagatagc	agtctgatca	180
cacatggtcc	aacaacactc	aaataataaa	tcaaataatna	tcagatgtta	aagattggtc	240
ttcaaacatc	atagccaatg	atgccccgct	tgcttataat	ctctccgaca	taaaaccaca	300
tcaacacctc	agtggccacc	aaaccattca	gcacagcttc	cttaactgtg	agctgtttga	360
agctaccagt	ctgagcacta	ttgactatnt	ttttcangct	ctgaatagct	ctagggatct	420
cagcangggg	gggaggaacc	agctcaacct	tggcggtant			459

&lt;210&gt; 143

&lt;211&gt; 140

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 143

acatttcctt	ccaccaagtc	aggactcctg	gcttctgtgg	gagttcttat	cacctgaggg	60
aaatccaaac	agtctctcct	agaaaggaat	agtgtcacca	acccccacca	tctccctgag	120
accatccgac	tccctgtgt					140

&lt;210&gt; 144

&lt;211&gt; 164

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(164)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 144

acttcagtaa	caacatacaa	taacaacatt	aagtgtatat	tgccatcttt	gtcattttct	60
atctatacca	ctctcccttc	tgaaaacaan	aatcactanc	caatcactta	tacaaatttg	120
aggcaattaa	tccatatttg	ttttcaataa	ggaaaaaaag	atgt		164

&lt;210&gt; 145

&lt;211&gt; 303

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(303)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 145

acgtagacca	tccaactttg	tatttgtaat	ggcaaacatc	cagnagcaat	tcctaaacaa	60
actggagggt	atttataccc	aattatccca	ttcattaaca	tgccctcttc	ctcaggctat	120
gcaggacagc	tatcataagt	cggcccaggc	atccagatac	taccatttgt	ataaacttca	180
gtaggggagt	ccatccaagt	gacaggtcta	atcaaaggag	gaaatggaac	ataagcccag	240
tagtaaaatn	ttgcttagct	gaaacagcca	caaaagactt	accgccgtgg	tgattaccat	300
caa						303

&lt;210&gt; 146

<211> 327  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(327)  
 <223> n = A,T,C or G

<400> 146  
 actgcagctc aattagaagt ggtctctgac ttctcatcanc ttctccctgg gctccatgac 60  
 actggcctgg agtgactcat tgctctgggtt ggttgagaga gctcctttgc caacaggcct 120  
 ccaagtccagg gctgggattt gtttcctttc cacattctag caacaatatg ctggccactt 180  
 cctgaacagg gaggggtggga ggagccagca tggaacaagc tgccactttc taaagtagcc 240  
 agacttgccc ctgggcctgt cacacctact gatgaccttc tgtgcttgca ggatggaatg 300  
 taggggtgag ctgtgtgact ctatgggt 327

<210> 147  
 <211> 173  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(173)  
 <223> n = A,T,C or G

<400> 147  
 acattgtttt tttagataa agcattgana gagctctcct taacgtgaca caatggaagg 60  
 actggaacac ataccacat ctttgttctg agggataatt ttctgataaa gtcttgctgt 120  
 atattcaagc acatatgtta tatattattc agttccatgt ttatagccta gtt 173

<210> 148  
 <211> 477  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(477)  
 <223> n = A,T,C or G

<400> 148  
 acaaccactt tatctcatcg aatttttaac ccaaactcac tcaactgtgcc ttcttatcct 60  
 atgggatata ttatttgatg ctccatttca tcacacatat atgaataata cactcatact 120  
 gccctactac ctgctgcaat aatcacattc ccttcctgtc ctgacctga agccattggg 180  
 gtggtcctag tggccatcag tccangcctg caccttgagc ccttgagctc cattgctcac 240  
 nccanccac ctcaccgacc ccactcctt acacagctac ctcttgctc tctaacccca 300  
 tagattatnt ccaaattcag tcaattaagt tactattaac actctaccg acatgtccag 360  
 caccactggg aagccttctc cagccaacac acacacacac acacncacac acacacatat 420  
 ccaggcacag gctacctcat cttcacaatc acccctttaa ttaccatgct atgggtgg 477

<210> 149  
 <211> 207  
 <212> DNA

```

<213> Homo sapien

<400> 149
acagttgtat tataatatca agaaataaac ttgcaatgag agcattttaag agggaagaac      60
taacgtatatt tagagagcca aggaagggtt ctgtgggggag tgggatgtaa ggtggggcct      120
gatgataaat aagagtcagc caggtaagtg ggtggtgtgg tatgggcaca gtgaagaaca      180
tttcaggcag agggaacagc agtgaag                                     207

<210> 150
<211> 111
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1) ... (111)
<223> n = A,T,C or G

<400> 150
accttgattt cattgctgct ctgatggaaa cccaactatc taatttagct aaaacatggg      60
cacttaaatg tggtcagtgt ttggacttgt taactantgg catctttggg t                111

<210> 151
<211> 196
<212> DNA
<213> Homo sapien

<400> 151
agcgcggcag gtcattattga acattccaga tacctatcat tactcgatgc tgttgataac      60
agcaagatgg ctttgaactc agggtcacca ccagctattg gaccttacta tgaaaaccat      120
ggataccaac cggaaaaccc ctatcccga cagcccactg tggccccac tgtctacgag      180
gtgcatccgg ctcatg                                     196

<210> 152
<211> 132
<212> DNA
<213> Homo sapien

<400> 152
acagcacttt cacatgtaag aaggagagaaa ttcctaaatg taggagaaa ataacagaac      60
cttccccctt tcatctagtg gtggaaacct gatgctttat gttgacagga atagaaccag      120
gaggagttt gt                                           132

<210> 153
<211> 285
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1) ... (285)
<223> n = A,T,C or G

<400> 153
acaanacca naganaggcca ctggccgtgg tgtcatggcc tccaaacatg aaagtgtcag      60

```

```

cttctgctct tatgtcctca tctgacaact ctttaccatt tttatcctcg ctcagcagga      120
gcacatcaat aaagtccaaa gtcttggact tggccttggc ttggaggaag tcatcaacac      180
cctggctagt gaggggtgcg cgccgtcctt ggatgacggc atctgtgaag tcgtgcacca      240
gtctgcaggc cctgtggaag cgccgtccac acggagtnag gaatt                        285

```

&lt;210&gt; 154

&lt;211&gt; 333

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 154

```

accacagtcc tgttggggcca gggcttcatg accctttctg tgaaaagcca tattatcacc      60
accccaaatt tttccttaaa tatctttaac tgaaggggtc agcctcttga ctgcaaagac      120
cctaagccgg ttacacagct aactcccact ggccctgatt tgtgaaattg ctgctgcctg      180
attggcacag gagtcgaagg tggtcagctc cctctctcgg tggaaacgaga ctctgatttg      240
agtttcacaa attctcgggc cacctcgtca ttgctcctct gaaataaaat ccggagaatg      300
gtcaggcctg tctcatccat atggatcttc cgg                                           333

```

&lt;210&gt; 155

&lt;211&gt; 308

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (308)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 155

```

actggaaata ataaaaccca catcacagtg ttgtgtcaaa gatcatcagg gcatggatgg      60
gaaagtgctt tgggaactgt aaagtgccta acacatgac gatgattttt gttataatat      120
ttgaatcacg gtgcatacaa actctcctgc ctgctcctcc tgggccccag cccagcccc      180
atcacagctc actgctctgt tcatccaggc ccagcatgta gtggctgatt cttcttggct      240
gcttttagcc tccanaagtt tctctgaagc caaccaaacc tctangtgta aggcatgctg      300
gccctggg                                           308

```

&lt;210&gt; 156

&lt;211&gt; 295

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 156

```

accttgctcg gtgcttggaa catattagga actcaaaata tgagatgata acagtgccta      60
ttattgatta ctgagagAAC tgtagacat ttagtgaag attttctaca caggaactga      120
gaataggaga ttatgttttg cctcatatt ctctcctatc ctcttgcct cattctatgt      180
ctaatatatt ctcaatcaaa taaggtttag ataatcagga aatcgaccaa ataccaatat      240
aaaaccagat gtctatcctt aagattttca aatagaaaac aaattaacag actat          295

```

&lt;210&gt; 157

&lt;211&gt; 126

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 157

```

acaagtttaa atagtgtgt cactgtgcat gtgctgaaat gtgaaatcca ccacatttct      60

```

gaagagcaaa acaaattctg tcatgtaatc tctatcttgg gtcgtgggta tatctgtccc 120  
cttagt 126

<210> 158  
<211> 442  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(442)  
<223> n = A,T,C or G

<400> 158  
accactgggt cttggaaaca cccatcctta atacgatgat ttttctgtcg tgtgaaaatg 60  
aanccagcag gctgccccta gtcagtcctt ccttcagag aaaaagagat ttgagaaagt 120  
gcctgggtaa ttcaccatta atttcctccc ccaaactctc tgagtcttcc cttaatattt 180  
ctggtgggtc tgaccaaagc aggtcatggg ttggttagca tttgggatcc cagtgaagta 240  
natgtttgta gccttgcata cttagccctt cccacgcaca aacggagtgg cagagtggg 300  
ccaaccctgt tttcccagtc cacgtagaca gattcacagt gcggaattct ggaagctgga 360  
nacagacggg ctctttgcag agccgggact ctgagangga catgagggcc tctgcctctg 420  
tgttcattct ctgatgtect gt 442

<210> 159  
<211> 498  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(498)  
<223> n = A,T,C or G

<400> 159  
acttccaggt aacgttggtg tttccggtga gcctgaactg atgggtgacg ttgtagggtc 60  
tccaacaaga actgaggttg cagagcgggt agggagagat gctgttccag ttgcacctgg 120  
gctgctgtgg actgttggtg attcctcact acggcccaag gttgtggaac tggcanaaag 180  
gtgtgtgtgt gganttgagc tcgggcgggt gtggtagggt gtgggtctt caacaggggc 240  
tgctgtgtgt ccgggagtg aangtggtgt gtcacttgag cttggccagc tctggaaagt 300  
antanattct tctgaaggc cagcgcttgt ggagctggca ngggtcantg ttgtgtgtaa 360  
cgaaccagtg ctgctgtggg tgggtgtana tcctccacaa agcctgaagt tatggtgtcn 420  
tcaggtaana atgtggttcc agtgcctctg ggcngctgtg gaaggttgta nattgtcacc 480  
aagggaataa gctgtggg 498

<210> 160  
<211> 380  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(380)  
<223> n = A,T,C or G

<400> 160

acctgcatcc agcttccctg ccaaactcac aaggagacat caacctctag acagggaaac	60
agcttcagga tacttccagg agacagagcc accagcagca aaacaaatat tcccatgcct	120
ggagcatggc atagaggaag ctganaaatg tggggctctga ggaagccatt tgagtctggc	180
cactagacat ctcatcagcc acttggtgtga agagatgccc catgacccca gatgcctctc	240
ccacccttac ctccatctca cacacttgag ctttccactc tgtataattc taacatcctg	300
gagaaaaatg gcagtttgac cgaacctgtt cacaacggta gaggctgatt tctaacgaaa	360
cttgtagaat gaagcctgga	380

&lt;210&gt; 161

&lt;211&gt; 114

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 161

actccacatc ccctctgagc aggcgggtgt cgttcaaggt gtatttggcc ttgcctgtca	60
cactgtccac tggccctta tccacttggt gcttaatccc tcgaaagagc atgt	114

&lt;210&gt; 162

&lt;211&gt; 177

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 162

actttctgaa tcgaatcaaa tgatacttag tgtagtttta atatcctcat atatatcaaa	60
gttttactac tctgataatt ttgtaaacca ggtaaccaga acatccagtc atacagcttt	120
tggtgatata taacttggca ataaccagc ctggtgatata ataaaactac tcactgt	177

&lt;210&gt; 163

&lt;211&gt; 137

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (137)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 163

catttataca gacaggcgtg aagacattca cgacaaaaac gcgaaattct atcccgtgac	60
canagaaggc agctacggct actcctacat cctggcgtgg gtggccttcg cctgcacctt	120
catcagcggc atgatgt	137

&lt;210&gt; 164

&lt;211&gt; 469

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (469)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 164

cttatcacia tgaatgttct cctgggcagc gttgtgatct ttgccacctt cgtgacttta	60
tgcaatgcat catgctattt catacctaag gagggagttc caggagattc aaccaggaaa	120

```

tgcattggatc tcaaaggaaa caaacaccca ataaactcgg agtggcagac tgacaactgt      180
gagacatgca cttgctacga aacagaaatt tcatgttgca cccttggttc tacacctgtg      240
ggttatgaca aagacaactg ccaaagaatc ttcaagaagg aggactgcaa gtatatcgtg      300
gtggagaaga aggacccaaa aaagacctgt tctgtcagtg aatggataat ctaatgtgct      360
tctagtaggc acagggctcc caggccaggc ctcattctcc tctggcctct aatagtcaat      420
gatttgtgtag ccattgcctat cagtaaaaag atntttgagc aaacacttt      469

```

&lt;210&gt; 165

&lt;211&gt; 195

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(195)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 165

```

acagtttttt atanatatcg acattgccgg cacttggtgtt cagtttcata aagctgggtgg      60
atccgctgctg atccactatt ccttggttag agtaaaaatt attcttatag cccatgtccc      120
tgcaggccgc ccgcccgtag ttctcggtcc agtcgtcttg gcacacaggg tgccaggact      180
tcctctgaga tgagt                                     195

```

&lt;210&gt; 166

&lt;211&gt; 383

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(383)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 166

```

acatcttagt agtgtggcac atcagggggc catcaggggc acagtcactc atagcctcgc      60
cgaggtcgga gtccacacca ccggtgtagg tgtgctcaat cttgggcttg gcgcccacct      120
ttggagaagg gatattgctg acacacatgt ccacaaagcc tgtgaactcg ccaaagaatt      180
tttgagacc agcctgagca aggggcggat gttcagcttc agtcctctct tcgtcagggtg      240
gatgccaaac tcgtctangg tccgtgggaa gctgggtgcc acntcaccta caacctgggc      300
gangatctta taaagagggt ccnagataaa ctccacgaaa cttctctggtg agctgctagt      360
nggggccttt ttggtgaact ttc                                     383

```

&lt;210&gt; 167

&lt;211&gt; 247

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(247)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 167

```

acagagccag accttggcca taaatgaanc agagattaag actaaacccc aagtcganat      60
tggagcagaa actggagcaa gaagtgggccc tggggctgaa gtagagacca aggccactgc      120

```



tatanccata cacagagcca actctcaggc caaggcnatg gttggggcag anccagagac 180  
 tcaatctgan tccaaagtgg tggctggaac actggtcatg acanaggcag tgactctgac 240  
 tgangtc 247

<210> 168

<211> 273

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(273)

<223> n = A,T,C or G

<400> 168

acttctaagt tttctagaag tggaaggatt gtantcatcc tgaaaatggg tttacttcaa 60  
 aatccctcan ccttggtctt cactactgtc tatactgana gtgtcatggt tccacaaagg 120  
 gctgacacct gagcctgnat ttctactcat ccctgagaag ccctttccag taggggtggc 180  
 aattcccaac ttccttgcca caagcttccc aggcctttctc ccctggaaaa ctccagcttg 240  
 agtcccagat acactcatgg gctgccctgg gca 273

<210> 169

<211> 431

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(431)

<223> n = A,T,C or G

<400> 169

acagccttgg ctccccaaa ctccacagtc tcagtgcaga aagatcatct tccagcagtc 60  
 agctcagacc aggggtcaaag gatgtgacat caacagtttc tggtttcaga acaggttcta 120  
 ctactgtcaa atgacccccc atacttcctc aaaggctgtg gtaagttttg cacaggtgag 180  
 ggcagcagaa aggggggtant tactgatgga caccatcttc tctgtatact ccacactgac 240  
 cttgccatgg gcaaaggccc ctaccacaaa aacaatagga tcaactgctgg gcaccagctc 300  
 acgcacatca ctgacaaccg ggatggaaaa agaantgcca actttcatac atccaactgg 360  
 aaagtgatct gatactggat tcttaattac cttcaaaagc ttctgggggc catcagctgc 420  
 tcgaacactg a 431

<210> 170

<211> 266

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(266)

<223> n = A,T,C or G

<400> 170

acctgtgggc tgggctgtta tgcctgtgcc ggctgctgaa agggagttca gaggtggagc 60  
 tcaaggagct ctgcaggcat ttgccaanc ctctccanag canagggagc aacctacact 120  
 ccccgctaga aagacaccag attggagtcc tgggaggggg agttgggggtg ggcatttgat 180

gtatacttgt cacctgaatg aangagccag agaggaanga gacgaanatg anattggcct 240  
tcaaagctag ggggtctggca ggtgga 266

<210> 171  
<211> 1248  
<212> DNA  
<213> Homo sapien  
  
<220>  
<221> misc\_feature  
<222> (1)...(1248)  
<223> n = A,T,C or G

<400> 171  
ggcagccaaa tcataaacgg cgaggactgc agcccgcact cgcagccctg gcaggcggca 60  
ctgggtcatgg aaaacgaatt gttctgctcg ggcgtcctgg tgcattccgca gtgggtgctg 120  
tcagccgcac actgtttcca gaagtgaatg cagagctcct acaccatcgg gctgggcttg 180  
cacagtcttg aggccgacca agagccaggg agccagatgg tggaggccag cctctccgta 240  
cggcaccacag agtacaacag acccttgctc gctaacgacc tcatgtcat caagttggac 300  
gaatccgtgt ccgagctga caccatccgg agcatcagca ttgcttcgca gtgccctacc 360  
gcggggaact cttgectcgt ttctggctgg ggtctgctgg cgaacggcag aatgcctacc 420  
gtgctgcagt gcgtgaacgt gtcgggtggg tctgaggagg tctgcagtaa gctctatgac 480  
ccgctgtacc accccagcat gttctgcgcc ggcggagggg aagaccagaa ggactcctgc 540  
aacggtgact ctggggggcc cctgatctgc aacgggtact tgcagggcct tgtgtctttc 600  
ggaaaagccc cgtgtggcca agttggcgtg ccagggtgct acaccaacct ctgcaaattc 660  
actgagtggg tagagaaaac cgtccaggcc agttaactct ggggactggg aacccatgaa 720  
attgaccccc aaatacatcc tgcggaagga attcaggaat atctgttccc agcccctcct 780  
ccctcaggcc caggagtcca ggccccccag ccctcctccc tcaaaccaag ggtacagatc 840  
cccagcccct cctccctcag acccaggagt ccagaccccc cagcccctcc tccctcagac 900  
ccaggagtcc agcccctcct ccctcagacc caggagtcca gacccccccag cccctcctcc 960  
ctcagaccca ggggtccagg cccccaaccc ctccctccctc agactcagag gtccaagccc 1020  
ccaaccntc attccccaga cccagagggtc cagggtcccag cccctcntcc ctcagaccca 1080  
gcggtccaat gccacctaga ctntccctgt acacagtgcc cccttggtggc acgttgaccc 1140  
aaccttacca gttggttttt catTTTTngt ccctttcccc tagatccaga aataaagttt 1200  
aagagaagng caaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 1248

<210> 172  
<211> 159  
<212> PRT  
<213> Homo sapien  
  
<220>  
<221> VARIANT  
<222> (1)...(159)  
<223> Xaa = Any Amino Acid

<400> 172  
Met Val Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro  
1 5 10 15  
Leu Leu Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser  
20 25 30  
Glu Ser Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr  
35 40 45  
Ala Gly Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Gly  
50 55 60

Arg Met Pro Thr Val Leu Gln Cys Val Asn Val Ser Val Val Ser Glu  
 65 70 75 80  
 Glu Val Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe  
 85 90 95  
 Cys Ala Gly Gly Gly Gln Xaa Gln Xaa Asp Ser Cys Asn Gly Asp Ser  
 100 105 110  
 Gly Gly Pro Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe  
 115 120 125  
 Gly Lys Ala Pro Cys Gly Gln Val Gly Val Pro Gly Val Tyr Thr Asn  
 130 135 140  
 Leu Cys Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Ala Ser  
 145 150 155

<210> 173

<211> 1265

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(1265)

<223> n = A,T,C or G

<400> 173

```

ggcagcccgc actcgcagcc ctggcaggcg gcactgggtca tggaaaacga attgttctgc      60
tcgggcgtcc tggtgcatcc gcagtgggtg ctgtcagccg cacactgttt ccagaactcc      120
tacaccatcg ggctgggacct gcacagtctt gaggccgacc aagagccagg gagccagatg      180
gtggaggcca gectctccgt acggcaccca gactacaaca gaccttgct cgctaacgac      240
ctcatgctca tcaagttgga cgaatccgtg tccgagtctg acaccatccg gagcatcagc      300
attgcttcgc agtgccttac cgcggggaac tcttgctctg tttctggctg gggctctgctg      360
gcgaacgggtg agctcacggg tgtgtgtctg ccctcttcaa ggaggctctc tgcccagtcg      420
cggggggtga cccagagctc tgcgtcccag gcagaatgcc taccgtgctg cagtgcgtga      480
acgtgtcggg ggtgtctgag gaggtctgca gtaagctcta tgaccgctg taccacccca      540
gcatgttctg cgccggcgga gggcaagacc agaaggactc ctgcaacggg gactctgggg      600
ggcccctgat ctgcaacggg tacttgacag gccttgtgtc tttcggaaaa gcccctgtg      660
gccaagttgg cgtgccaggt gtctacacca acctctgcaa attcactgag tggatagaga      720
aaaccgtcca ggccagttaa ctctggggac tgggaaccca tgaaattgac ccccaatac      780
atcctgcgga aggaattcag gaatatctgt tcccagcccc tcctccctca ggcccaggag      840
tccaggcccc cagccctctc tccctcaaac caaggttaca gatccccagc ccctcctccc      900
tcagaccagc gagtccagac cccccagccc ctctccctc agaccagga gtccagcccc      960
tcctccntca gaccaggag tccagacccc ccagcccctc ctccctcaga cccaggggtt     1020
gaggccccca accctctctc ctccagagtc agaggtccaa gcccacaacc cctcgttccc     1080
cagaccagga ggttnaggtc ccagcccctc ttcntcaga cccagnggtc caatgccacc     1140
tagattttcc ctgnacacag tgcccccttg tggngangttg acccaacctt accagttggt     1200
ttttcatttt tngtcccttt cccctagatc cagaaataaa gtttaagaga ngngcaaaaa     1260
aaaaa                                           1265

```

<210> 174

<211> 1459

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(1459)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 174

ggtcagccgc	acactgtttc	cagaagtgag	tgcagagctc	ctacaccatc	gggctgggcc	60
tgcacagtct	tgaggccgac	caagagccag	ggagccagat	ggaggaggcc	agcctctccg	120
tacggcacc	agagtacaac	agacccttgc	tcgctaacga	cctcatgctc	atcaagttgg	180
acgaatccgt	gtccgagctc	gacaccatcc	ggagcatcag	cattgcttcg	cagtgcctta	240
ccgcggggaa	ctcttgccct	gtttctggct	ggggtctgct	ggcgaacggg	gagctcacgg	300
gtgtgtgtct	gccctcttca	aggaggtcct	ctgcccagtc	gcgggggctg	acccagagct	360
ctgctgcccc	ggcagaatgc	ctaccgtgct	gcagtgcgtg	aacgtgtcgg	tggtgtctga	420
ngaggctctg	antaagctct	atgaccgct	gtaccacccc	ancatgttct	gcgccggcgg	480
agggcaagac	cagaaggact	cctgcaacgt	gagagagggg	aaaggggagg	gcaggcgact	540
cagggaaagg	tggagaaggg	ggagacagag	acacacaggg	ccgcatggcg	agatgcagag	600
atggagagac	acacagggag	acagtgacaa	ctagagagag	aaactgagag	aaacagagaa	660
ataaacacag	gaataaagag	aagcaaagga	agagagaaac	agaaacagac	atggggaggc	720
agaaacacac	acacatagaa	atgcagttga	ccttccaaca	gcatggggcc	tgagggcggt	780
gacctccacc	caatagaaaa	tcctcttata	acttttgact	ccccaaaaac	ctgactagaa	840
atagcctact	gttgacgggg	agccttacca	ataacataaa	tagtcgattt	atgcatacgt	900
tttatgcatt	catgatatac	ctttgttgga	atTTTTtgat	atttctaagc	tacacagttc	960
gtctgtgaat	TTTTTTaaat	tgttgcaact	ctcctaaaaa	TTTTctgatg	tgtttattga	1020
aaaaatccaa	gtataagtgg	acttgtgcat	tcaaaccagg	gttgttcaag	ggtcaactgt	1080
gtacccagag	ggaaacagtg	acacagattc	atagaggtga	aacacgaaga	gaaacaggaa	1140
aaatcaagac	tctacaaaga	ggctgggcag	ggagggtcat	gcctgtaatc	ccagcacttt	1200
gggaggcgag	gcaggcgag	cacttgaggt	aaggagttca	agaccagcct	ggccaaaatg	1260
gtgaaatcct	gtctgtacta	aaaatacaaa	agttagctgg	atatggtggc	aggcgctgt	1320
aatcccgct	acttgggagg	ctgaggcagg	agaattgctt	gaatatggga	ggcagaggtt	1380
gaagtgaagt	gagatcacac	cactatactc	cagctggggc	aacagagtaa	gactctgtct	1440
caaaaaaaaa	aaaaaaaaa					1459

&lt;210&gt; 175

&lt;211&gt; 1167

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (1167)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 175

gcgcagccct	ggcaggcggc	actgggtcatg	gaaaacgaat	tgttctgctc	gggcgtcctg	60
gtgcatccgc	agtgggtgct	gtcagccgca	cactgtttcc	agaactccta	caccatcggg	120
ctgggcctgc	acagtcttga	ggccgaccaa	gagccaggga	gccagatggg	ggaggccagc	180
ctctccgtac	ggcaccaga	gtacaacaga	ctcttgctcg	ctaacgacct	catgctcatc	240
aagttggacg	aatccgtgtc	cgagtctgac	accatccgga	gcacagcat	tgcttcgcag	300
tgcctaccg	cggggaactc	ttgcctcgtn	tctggctggg	gtctgctggc	gaacggcaga	360
atgcctaccg	tgctgcactg	cgtgaacgtg	tgggtgggtg	ctgaggangt	ctgcagtaag	420
ctctatgacc	cgctgtacca	ccccagcatg	ttctgcgccg	gcggagggca	agaccagaag	480
gactcctgca	acggtgactc	tgggggggcc	ctgatctgca	acgggtactt	gcagggcctt	540
gtgtctttcg	gaaaagcccc	gtgtggccaa	cttggcgtgc	caggtgtcta	caccaacctc	600
tgcaaattca	ctgagtggat	agagaaaacc	gtccagncca	gttaactctg	gggactggga	660
acccatgaaa	ttgacccccca	aatacatcct	gcggaangaa	ttcaggaata	tctgttccca	720
gccccctctc	cctcaggccc	aggagtccag	gccccagcc	cctcctccct	caaaccaagg	780
gtacagatcc	ccagcccctc	ctccctcaga	cccaggagtc	cagaccccc	agccccctnt	840
cctcagacc	caggagtcca	gccccctctc	cntcagacgc	aggagtccag	accccccagc	900

```

ccntcntccg tcagacccag ggggtgcaggc ccccaacccc tcntccntca gagtcagagg 960
tccaagcccc caaccctcgc ttccccagac ccagaggtnc aggtcccagc cctcctcccc 1020
tcagacccag cgggtccaatg ccacctagan tntccctgta cacagtgcgc ccttggtggca 1080
ngttgaccca accttaccag ttgggtttttc attttttgtc cctttcccct agatccagaa 1140
ataaagtnta agagaagcgc aaaaaaa 1167

```

&lt;210&gt; 176

&lt;211&gt; 205

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; VARIANT

&lt;222&gt; (1)...(205)

&lt;223&gt; Xaa = Any Amino Acid

&lt;400&gt; 176

```

Met Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln Trp
1          5          10          15
Val Leu Ser Ala Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu
20          25          30
Gly Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val
35          40          45
Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Leu Leu Leu
50          55          60
Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu Ser
65          70          75          80
Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala Gly
85          90          95
Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Gly Arg Met
100          105          110
Pro Thr Val Leu His Cys Val Asn Val Ser Val Val Ser Glu Xaa Val
115          120          125
Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe Cys Ala
130          135          140
Gly Gly Gly Gln Asp Gln Lys Asp Ser Cys Asn Gly Asp Ser Gly Gly
145          150          155          160
Pro Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe Gly Lys
165          170          175
Ala Pro Cys Gly Gln Leu Gly Val Pro Gly Val Tyr Thr Asn Leu Cys
180          185          190
Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Xaa Ser
195          200          205

```

&lt;210&gt; 177

&lt;211&gt; 1119

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 177

```

gcgcactcgc agccctggca ggcggcactg gtcattggaaa acgaattggt ctgctcgggc 60
gtcctgggtgc atccgcagtg ggtgctgtca gccgcacact gtttcagaaa ctctacacc 120
atcgggcttg gctgcacag tcttgaggcc gaccaagagc cagggagcca gatgggtggag 180
gccagcctct ccgtacggca cccagagtac aacagaccct tgctcgctaa cgacctcatg 240
ctcatcaagt tggacgaatc cgtgtccgag tctgacacca tccggagcat cagcattgct 300

```

```

tcgcagtgcc ctaccgcggg gaactcttgc ctcgtttctg gctgggggtct gctggcgaaac 360
gatgctgtga ttgccatcca gtcccagact gtgggaggct gggagtgatga gaagctttcc 420
caaccctggc aggggtgtac catttcggca acttccagtg caaggacgtc ctgctgcatc 480
ctcactgggt gctcactact gctcactgca tcaccgcgaa cactgtgatc aactagccag 540
caccatagtt ctccgaagtc agactatcat gattactgtg ttgactgtgc tgtctattgt 600
actaaccatg ccgatgttta ggtgaaatta gcgtcacttg gcctcaacca tcttggtatc 660
cagttatcct cactgaattg agatttcctg cttcagtgtc agccattccc acataatttc 720
tgacctacag aggtgagggg tcatatagct cttcaaggat gctgggtactc ccttcacaaa 780
ttcattttctc ctgttgtagt gaaagggtgc cctctggag cctcccaggg tgggtgtgca 840
ggtcacaatg atgaatgtat gatcgtgttc ccattaccca aagcctttaa atccctcatg 900
ctcagtacac cagggcaggt ctagcatttc ttcathtagt gtatgctgtc cattcatgca 960
accacctcag gactcctgga ttctctgcct agttgagctc ctgcatgctg cctccttggg 1020
gaggtgaggg agagggccca tggttcaatg ggatctgtgc agttgtaaca cattaggtgc 1080
ttaataaaca gaagctgtga tgttaaaaaa aaaaaaaaaa 1119

```

&lt;210&gt; 178

&lt;211&gt; 164

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; VARIANT

&lt;222&gt; (1)...(164)

&lt;223&gt; Xaa = Any Amino Acid

&lt;400&gt; 178

```

Met Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln Trp
 1          5          10          15
Val Leu Ser Ala Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu
          20          25          30
Gly Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val
          35          40          45
Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro Leu Leu
          50          55          60
Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu Ser
65          70          75          80
Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala Gly
          85          90          95
Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Asp Ala Val
          100          105          110
Ile Ala Ile Gln Ser Xaa Thr Val Gly Gly Trp Glu Cys Glu Lys Leu
          115          120          125
Ser Gln Pro Trp Gln Gly Cys Thr Ile Ser Ala Thr Ser Ser Ala Arg
          130          135          140
Thr Ser Cys Cys Ile Leu Thr Gly Cys Ser Leu Leu Leu Thr Ala Ser
145          150          155          160
Pro Gly Thr Leu

```

&lt;210&gt; 179

&lt;211&gt; 250

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 179

```

ctggagtgcc ttggtgtttc aagcccctgc aggaagcaga atgcaccttc tgaggcacct      60
ccagctgccc ccggccgggg gatgcgaggc tcggagcacc cttgcccggc tgtgattgct      120
gccaggcact gttcatctca gcttttctgt ccctttgctc ccggcaagcg cttctgctga      180
aagttcatat ctggagcctg atgtcttaac gaataaaggt cccatgctcc acccgaaaaa      240
aaaaaaaaaa                                     250

```

&lt;210&gt; 180

&lt;211&gt; 202

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 180

```

actagtccag tgtggtggaa ttccattgtg ttggggcccaa cacaatggct acctttaaca      60
tcacccagac cccgcccctg cccgtgcccc acgtgctgct taacgacagt atgatgctta      120
ctctgctact cggaaactat ttttatgtaa ttaatgtatg ctttcttggt tataaatgcc      180
tgattttaaa aaaaaaaaaa aa                                     202

```

&lt;210&gt; 181

&lt;211&gt; 558

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (558)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 181

```

tccytttght naggtttkkg agacamccck agacctwaan ctgtgtcaca gacttcynng      60
aatgtttagg cagtgttagt aatttcytcg taatgattct gttattactt tcctnattct      120
ttattcctct ttcttctgaa gattaatgaa gttgaaaatt gaggtggata aatacaaaaa      180
ggtagtgtga tagtataagt atctaagtcg agatgaaagt gtgttatata tatccattca      240
aaattatgca agttagtaat tactcagggg taactaaatt actttaatat gctgttgaac      300
ctactctgtt ccttggctag aaaaaattat aaacaggact ttgttagttt gggaagccaa      360
attgataata ttctatgttc taaaagttgg gctatacata aattattaag aaatatggaw      420
ttttattccc aggaatatgg kgttcatttt atgaatatta cscrggatag awgtwtgagt      480
aaaaycagtt ttggtwaata ygtwaatatg tcmtaaataa acaakgcttt gacttatttc      540
caaaaaaaaa aaaaaaaaaa                                     558

```

&lt;210&gt; 182

&lt;211&gt; 479

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (479)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 182

```

acagggwttk grggatgcta agsccccrga rwtggtttga tccaaccctg gcttwttttc      60
agaggggaaa atggggccta gaagttacag mscatytagy tgggtgcgmg gcacccctgg      120
cstcacacag astcccgagt agctgggact acaggcacac agtcaactgaa gcaggccctg      180
ttwgcaattc acgttgccac ctccaactta aacattcttc atatgtgatg tccttagtca      240
ctaaggttaa actttcccac ccagaaaagg caacttagat aaaatcttag agtactttca      300

```

```
tactmttcta agtcctcttc cagcctcact kkgagtcctm cytggggggt gataggaant 360
ntctcttggc tttctcaata aartctctat ycatctcatg ttaatttgg tacgcatara 420
awtgstgara aaattaaaat gttctgggty mactttaaaa araaaaaaaa aaaaaaaaaa 479
```

&lt;210&gt; 183

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 183

```
aggcgggagc agaagctaaa gccaaagccc aagaagagtg gcagtgccag cactgggtgcc 60
agtaccagta ccaataacag tgccagtgcc agtgccagca ccagtgggtg cttcagtgtc 120
ggtgccagcc tgaccgccac tctcacattt gggctcttcg ctggccttgg tggagctggg 180
gccagcacca gtggcagctc tgggtgcctgt ggtttctcct acaagtgaga ttttagatat 240
tgtaatcct gccagtcttt ctcttcaagc caggggtgcat cctcagaaac ctactcaaca 300
cagcactcta ggcagccact atcaatcaat tgaagttgac actctgcatt aratctattt 360
gccatttcaa aaaaaaaaaa aaaa 384
```

&lt;210&gt; 184

&lt;211&gt; 496

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(496)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 184

```
accgaattgg gaccgctggc ttataagcga tcatgtyynt ccrgtatkac ctcaacgagc 60
agggagatcg agtctatacg ctgaagaaat ttgacccgat gggacaacag acctgtctag 120
cccactctgc tcggttctcc ccagatgaca aatactctsg acaccgaatc accatcaaga 180
aacgcttcaa ggtgctcatg accagcaac cgcgcctgt cctctgaggg tcccttaaac 240
tgatgtcttt tctgccacct gttacccctc ggagactccg taaccaaact cttcggaactg 300
tgagccctga tgcctttttg ccagccatac tctttggcat ccagtctctc gtggcgattg 360
attatgcttg tgtgaggcaa tcatgggtggc atcaccata aagggaacac atttgacttt 420
tttttctcat attttaaatt actacmagaw tattwmagaw waaatgawtt gaaaaactst 480
taaaaaaaaa aaaaaa 496
```

&lt;210&gt; 185

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 185

```
gctggtagcc tatggcgkgg cccacggagg ggctcctgag gccacggrac agtgacttcc 60
caagtatcyt gcgcsgcgtc ttctaccgtc cctacctgca gatcttcggg cagattcccc 120
aggaggacat ggacgtggcc ctcatggagc acagcaactg ytcgtcggag cccggcttct 180
gggcacaccc tcctggggcc caggcgggca cctgcgtctc ccagtatgcc aactggctgg 240
tggtgctgct cctcgtcatc ttctgctcgt tggccaacat cctgctgggtc aacttgctca 300
ttgccatgtt cagttacaca ttcggaacag tacagggcaa cagegatctc tactgggaag 360
gcgcagcgtt accgctcat ccgg 384
```

&lt;210&gt; 186

&lt;211&gt; 577



&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (577)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 186

gagttagctc	ctccacaacc	ttgatgaggt	cgtctgcagt	ggcctctcgc	ttcataccgc	60
tnccatcgtc	atactgtagg	tttgccacca	cytcctggca	tcttggggcg	gcntaatatt	120
ccaggaaact	ctcaatcaag	tcaccgtcga	tgaaacctgt	gggctgggtc	tgtcttccgc	180
tcgggtgtgaa	aggatctccc	agaaggagtg	ctcgatcttc	cccacacttt	tgatgacttt	240
attgagtcga	ttctgcatgt	ccagcaggag	gttgtaggag	ctctctgaca	gtgaggtcac	300
cagccctatc	atgccgttga	mcgtgccgaa	garcaccgag	ccttgtgtgg	gggkkgaggt	360
ctcaccacaga	ttctgcatta	ccagagagcc	gtggcaaaag	acattgacaa	actcgccag	420
gtggaaaaag	amcamctcct	ggargtgctn	gccgtcctc	gtcmgttggt	ggcagcgctw	480
tccttttgac	acacaaacaa	gttaaaggca	ttttcagccc	ccagaaantt	gtcatcatcc	540
aagatntcgc	acagcactna	tccagttggg	attaaat			577

&lt;210&gt; 187

&lt;211&gt; 534

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (534)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 187

aacatcttcc	tgtataatgc	tgtgtaatat	cgatccgatn	ttgtctgstg	agaatycatw	60
actkggaaaa	gmaacattaa	agcctggaca	ctgggtattaa	aattcacaa	atgcaacact	120
ttaaacagtg	tgtcaatctg	ctcccyynac	tttgtcatca	ccagtctggg	aakaagggtg	180
tgccctattc	acacctgtta	aaagggcgct	aagcattttt	gattcaacat	cttttttttt	240
gacacaagtc	cgaaaaaagc	aaaagtaaac	agttatyaat	ttgttagcca	attcactttc	300
ttcatgggac	agagccatyt	gatttaaaaa	gcaaattgca	taatattgag	cttygggagc	360
tgatatttga	gcggaagagt	agcctttcta	cttcaccaga	cacaactccc	tttcatattg	420
ggatgttnac	naaagtwatg	tctctwacag	atgggatgct	tttgtggcaa	ttctgttctg	480
aggatctccc	agtttattta	ccacttgcac	aagaaggcgt	tttcttctc	aggc	534

&lt;210&gt; 188

&lt;211&gt; 761

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (761)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 188

agaaaccagt	atctctnaaa	acaacctctc	ataccttggtg	gacctaat	ttgtgtgcgtg	60
tgtgtgtgcg	cgcatattat	atagacaggc	acatcttttt	tacttttgta	aaagcttatg	120
cctctttggg	atctatatct	gtgaaagttt	taatgatctg	ccataatgtc	ttggggacct	180

ttgtcttctg	tgtaaattggt	actagagaaa	acacctatnt	tatgagtcaa	tctagttngt	240
tttattcgac	atgaaggaaa	tttccagatn	acaacactna	caaactctcc	ctkgackarg	300
ggggacaaag	aaaagcaaaa	ctgamcataa	raaacaatwa	cctgggtgaga	arttgcataa	360
acagaaatwr	ggtagtatat	tgaarnacag	catcattaaa	rmgtttwtktt	wtctctccctt	420
gcaaaaaaca	tgtacngact	tcccgttgag	taatgccaag	ttgttttttt	tatnataaaa	480
cttgcccttc	attacatggt	tnaaagtggg	gtgggtgggcc	aaaatattga	aatgatggaa	540
ctgactgata	aagctgtaca	aataagcagt	gtgcctaaca	agcaacacag	taatgttgac	600
atgcttaatt	cacaaatgct	aatttcatta	taaatgtttg	ctaaaataca	ctttgaacta	660
tttttctgtn	ttcccagagc	tgagatntta	gattttatgt	agtatnaagt	gaaaaantac	720
gaaaataata	acattgaaga	aaaananaaa	aaanaaaaaa	a		761

&lt;210&gt; 189

&lt;211&gt; 482

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (482)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 189

tttttttttt	tttgccgatn	ctactatttt	attgcaggan	gtgggggtgt	atgcaccgca	60
caccggggct	atnagaagca	agaaggaagg	agggagggca	cagccccttg	ctgagcaaca	120
aagccgcctg	ctgccttctc	tgtctgtctc	ctgggtgcagg	cacatgggga	gaccttcccc	180
aaggcagggg	ccaccagtc	aggggtggga	atacaggggg	tgggangtgt	gcataagaag	240
tgataggcac	aggccacccg	gtacagaccc	ctcggctcct	gacaggtnga	tttcgaccag	300
gtcattgtgc	cctgcccagg	cacagcgtn	atctggaaaa	gacagaatgc	tttccttttc	360
aaatttggct	ngtcatngaa	ngggcanttt	tccaanttn	gctnggtcct	ggtacncttg	420
gttcggccca	gtccnccgtc	caaaaantat	tcaccnct	ccnaattgct	tgcnggnccc	480
cc						482

&lt;210&gt; 190

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (471)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 190

tttttttttt	ttttaaaaca	gtttttcaca	acaaaattta	ttagaagaat	agtggttttg	60
aaaactctcg	catccagtga	gaactaccat	acaccacatt	acagctngga	atgtnctcca	120
aatgtctggg	caaatgatag	aatggaacca	ttcaatctta	cacatgcacg	aaagaacaag	180
cgcttttgac	atacaatgca	caaaaaaaaa	aggggggggg	gaccacatgg	attaaaattt	240
taagtactca	tcacatacat	taagacacag	ttctagtcca	gtcnaaaatc	agaactgcnt	300
tgaaaaattt	catgtatgca	atccaaccaa	agaacttnat	tgggtgatcat	gantnctcta	360
ctacatcnac	cttgatcatt	gccaggaaacn	aaaagttnaa	ancacncngt	acaaaaanaa	420
tctgtaattn	anttcaacct	ccgtacngaa	aaatnttnt	tatacactcc	c	471

&lt;210&gt; 191

&lt;211&gt; 402

&lt;212&gt; DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(402)

<223> n = A,T,C or G

<400> 191

gagggattga aggtctgttc tastgtcggm ctgttcagcc accaactcta acaagttgct	60
gtcttccact cactgtctgt aagcttttta acccagacwg tatcttcata aatagaacaa	120
attcttcacc agtcacatct tctaggacct ttttggattc agttagtata agctcttcca	180
cttcctttgt taagacttca tctggtaaag tcttaagttt tgtagaaagg aattyaattg	240
ctcgttctct aacaatgtcc tctccttgaa gtatttggtt gaacaaccca cctaaagtcc	300
ctttgtgcat ccattttaaa tatacttaat agggcattgk tncactaggt taaattctgc	360
aagagtcacg tgtctgcaaa agttgcgtta gtatatctgc ca	402

<210> 192

<211> 601

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(601)

<223> n = A,T,C or G

<400> 192

gagctcggat ccaataatct ttgtctgagg gcagcacaca tatncagtgc catggnaact	60
ggctctacccc acatgggagc agcatgccgt agntatataa ggctattccc tgagtcagac	120
atgcytyttt gaytaccgtg tgccaagtgc tgggtgattct yaacacacyt ccatcccgtt	180
cttttgtgga aaaactggca cttktctgga actagcarga catcacttac aaattcacc	240
acgagacact tgaaagggtg aacaaagcga ytcttgcat gctttttgtc cctccggcac	300
cagttgtcaa tactaaccgc ctggtttgcc tccatcacat ttgtgatctg tagctctgga	360
tacatctcct gacagtactg aagaacttct tcttttgttt caaaagcarg tcttggtgcc	420
tgttggatca gggtcccat tcccagtcyg aatgttcaca tggcatattt wacttcccac	480
aaaacattgc gatttgaggc tcagcaacag caaatcctgt tccggcattg gctgcaagag	540
cctcgatgta gccggccagc gccaaaggcag gcgccgtgag cccaccagc agcagaagca	600
g	601

<210> 193

<211> 608

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(608)

<223> n = A,T,C or G

<400> 193

atacagccca natcccacca cgaagatgag cttgttgact gagaacctga tgcggtcact	60
gggtcccgctg tagccccagc gactctccac ctgctggaag cggttgatgc tgcactcytt	120
cccaacgcag gcagmagcgg gscgggtcaa tgaactccay tcgtggcttg gggtkgacgg	180
tkaagtgcag gaagaggctg accacctgcg gggtccaccag gatgcccgac tgtgcgggac	240
ctgcagcgaa actcctcgat ggtcatgagc ggggaagcgaa tgaggcccag ggccttgccc	300

```

agaaccttcc gctgtttctc tggcgctcacc tgcagctgct gccgctgaca ctgggcctcg      360
gaccagcgga caaacggcrt tgaacagccg cacctcacgg atgcccagtg tgtcgcgctc      420
caggammgsc accagcgtgt ccaggtcaat gtcggtgaag ccctccgcgg gtrattggcgt      480
ctgcagtgtt tttgtcgatg ttctccaggc acaggctggc cagctgcggt tcatcgaaga      540
gtcgcgcctg cgtgagcagc atgaaggcgt tgtcggctcg cagttcttct tcaggaactc      600
cacgcaat                                         608

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&lt;210&gt; 194

&lt;211&gt; 392

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(392)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 194

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gaacggctgg accttgctc gcattgtgct tgctggcagg gaataccttg gcaagcagyt      60
ccagtcagag cagccccaga ccgctgccgc ccgaagctaa gcctgcctct ggccttcccc      120
tccgctcaa tgcagaacca gtagtgggag cactgtgttt agagttaaga gtgaacactg      180
tttgatttta cttgggaatt tctctgttta tatagctttt cccaatgcta atttccaaac      240
aacaacaaca aaataacatg tttgcctgtt aagttgtata aaagtaggtg attctgtatt      300
taaagaaaat attactgtta catatactgc ttgcaatttc tgtatttatt gktnctstgg      360
aaataaatat agttattaaa ggttgtcant cc                                         392

```

&lt;210&gt; 195

&lt;211&gt; 502

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(502)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 195

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ccsttkgagg ggtkaggkyc cagttyccga gtggaagaaa caggccagga gaagtgcgtg      60
ccgagctgag gcagatgttc ccacagtgc cccagagacc stgggstata gtytctgacc      120
cctcncaagg aaagaccacs ttctggggac atgggctgga gggcaggacc tagaggcacc      180
aaggggaagg cccattccgg ggstgttccc cgaggaggaa ggggaaggggc tctgtgtgcc      240
ccccasgagg aagaggccct gagtccctgg atcagacacc ctttcacgtg tatccccaca      300
caaatgcaag ctcaccaagg tcccctctca gtccccttcc stacaccctg amcggccact      360
gscscacacc caccagagc acgccaccg ccatggggar tgtgctcaag gartcgcnng      420
gcarcgtgga catctngtcc cagaaggggg cagaatctcc aatagangga ctgarcmstt      480
gctnanaaaa aaaaanaaaa aa                                         502

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&lt;210&gt; 196

&lt;211&gt; 665

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(665)

<223> n = A,T,C or G

<400> 196

ggttacttgg	tttcattgcc	accacttagt	ggatgtcatt	tagaaccatt	ttgtctgctc	60
cctctggaag	ccttgcgag	agcggacttt	gtaattgttg	gagaataact	gctgaatttt	120
wagctgtttk	gagttgatts	gcaccactgc	accacaact	tcaatatgaa	aacyawttga	180
actwatttat	tatcttgtga	aaagtataac	aatgaaaatt	ttgttcatac	tgtattkac	240
aagtatgatg	aaaagcaawa	gatatatatt	cttttattat	gttaaattat	gattgccatt	300
attaatcggc	aaaatgtgga	gtgtatgttc	ttttcacagt	aatatatgcc	ttttgtaact	360
tcacttggtt	attttattgt	aaatgarta	caaaattctt	aatttaagar	aatggatgt	420
watatttatt	tcattaattt	ctttcctkgt	ttacgtwaat	tttgaaaaga	wtgcatgatt	480
tcttgacaga	aatcgatctt	gatgctgtgg	aagtagtttg	accacatcc	ctatgagttt	540
ttcttagaat	gtataaagg	tgtagcccat	cnaacttcaa	agaaaaaat	gaccacatac	600
tttgaatca	ggctgaaatg	tggcatgctn	ttctaattcc	aactttataa	actagcaaan	660
aagtg						665

<210> 197

<211> 492

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1) ... (492)

<223> n = A,T,C or G

<400> 197

ttttnttttt	ttttttttgc	aggaaggatt	ccattttattg	tggatgcatt	ttcacaatat	60
atgtttattg	gagcgatcca	ttatcagtga	aaagtatcaa	gtgtttataa	natttttagg	120
aaggcagatt	cacagaacat	gctngtcngc	ttgcagtttt	acctcgtana	gatnacagag	180
aattatagtc	naaccagtaa	acnaggaatt	tacttttcaa	aagattaaat	ccaaactgaa	240
caaaattcta	ccctgaaact	tactccatcc	aaatattgga	ataanagtca	gcagtatac	300
attctcttct	gaactttaga	ttttctagaa	aaatatgtaa	tagtgatcag	gaagagctct	360
tgttcaaaaag	tacaacnaag	caatgttccc	ttaccatagg	ccttaattca	aactttgatc	420
catttcactc	ccatcacggg	agtcaatgct	acctgggaca	cttgtatttt	gttcatnctg	480
ancntggctt	aa					492

<210> 198

<211> 478

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1) ... (478)

<223> n = A,T,C or G

<400> 198

ttttnttttgn	atttcantct	gtannaanta	ttttcattat	gtttattana	aaaatatnaa	60
tgtntccacn	acaaatcatn	ttacntnagt	aagaggccan	ctacattgta	caacatacac	120
tgagtatatt	ttgaaaagga	caagtttaaa	gtanacncat	attgccganc	atancacatt	180
tatacatggc	ttgattgata	tttagcacag	canaaactga	gtgagttacc	agaaanaaat	240
natatatgtc	aatcngattt	aagatacaaa	acagatccta	tggtacatan	catcntgtag	300
gagttgtggc	tttatgttta	ctgaaagtca	atgcagttcc	tgtacaaaga	gatggccgta	360
agcattctag	tacctctact	ccatgggttaa	gaatcgtaca	cttatgttta	catatgtnc	420

gggtaagaat tgtgttaagt naanttatgg agagggtccan gagaaaaatt tgatncaa 478

<210> 199

<211> 482

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(482)

<223> n = A,T,C or G

<400> 199

agtgacttgt cctccaacaa aacccttga tcaagtttgt ggcactgaca atcagacct	60
tgctagtccc tgcacttat tgcactactaa atgcagactg gaggggacca aaaaggggca	120
tcaactccag ctggattatt ttggagcctg caaatctatt cctacttgta cggactttga	180
agtgattcag tttcctctac ggatgagaga ctgggtcaag aatatacctca tgcagcttta	240
tgaagccnac tctgaacacg ctggttatct nagatgagaa ncagagaaat aaagtcnaga	300
aaatttacct ggangaaaag aggctttngg ctggggacca tccattgaa ccttctctta	360
anggacttta agaanaaact accacatgtn tgtngtatcc tgggtgccngg ccgtttantg	420
aacntngacn ncacccttnt ggaatanant cttgacngcn tcctgaactt gtcctctgc	480
ga.	482

<210> 200

<211> 270

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(270)

<223> n = A,T,C or G

<400> 200

cggccgcaag tgcaactcca gctggggccg tgcggacgaa gattctgcca gcagttggtc	60
cgactgcgac gacggcgccg gcgacagtcg cagggtgcagc gcgggcccct ggggtcttgc	120
aaggctgagc tgacgccgca gaggtcgtgt cacgtcccac gaccttgacg ccgtcgggga	180
cagccggaac agagcccggg gaangcggga ggcctcgggg agcccctcgg gaagggcggc	240
ccgagagata cgcaggtgca ggtggccgcc	270

<210> 201

<211> 419

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(419)

<223> n = A,T,C or G

<400> 201

tttttttttt ttttggaaac tactgcgagc acagcaggtc agcaacaagt ttattttgca	60
gctagcaagg taacagggtta gggcatgggt acatgttcag gtcaacttcc tttgtcgtgg	120
ttgattgggt tgtctttatg ggggccccgg ggggtagggg aaanccaagc anaantaaca	180
tggagtgggt gcaccctccc tgtagaacct gggtacnaaa gcttggggca gttcacctgg	240

tctgtgaccg	tcattttctt	gacatcaatg	ttattagaag	tcaggatata	ttttagagag	300
tccactgtnt	ctggagggag	attaggggtt	cttgccaana	tccaancaaa	atccacntga	360
aaaagttgga	tgatncangt	acngaatacc	ganggcatan	ttctcatant	cggtggcca	419

<210> 202  
 <211> 509  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(509)  
 <223> n = A,T,C or G

<400> 202						
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tggcacttaa	tccattttta	tttcaaaatg	tctacaaant	ttnaatncnc	cattatacng	120
gtnattttnc	aaaatctaaa	nntttattca	atntnagcca	aantccttac	ncaaatnnaa	180
tacnncnaaa	aatcaaaaaat	atacntntct	ttcagcaaac	ttngttacat	aaattaaaaa	240
aatatatacg	gctgggtgtt	tcaaagtaca	attatcttaa	cactgcaaac	atnttttnaa	300
ggaactaaaa	taaaaaaaaa	cactnccgca	aagggttaaag	ggaacaacaa	attcntttta	360
caacancnnc	nattataaaa	atcataatct	aaatcttagg	ggaatatata	cttcacacng	420
ggatcttaac	ttttactnca	ctttgtttat	ttttttanaa	ccattgtntt	gggccaaca	480
caatggnaat	ncnccnncnc	tggtactagt				509

<210> 203  
 <211> 583  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(583)  
 <223> n = A,T,C or G

<400> 203						
tttttttttt	ttttttttga	ccccctctt	ataaaaaaca	agttaccatt	ttattttact	60
tacacatatt	tattttataa	ttggtattag	atattcaaaa	ggcagctttt	aaaatcaaac	120
taaatggaaa	ctgccttaga	tacataattc	ttaggaatta	gcttaaaatc	tgccataaagt	180
gaaaatcttc	tctagctctt	ttgactgtaa	atttttgact	cttgtaaaac	atccaaattc	240
atttttcttg	tctttaaaat	tatctaattc	ttccattttt	tccctattcc	aagtcaattt	300
gcttctctag	cctcatttcc	tagctcttat	ctactattag	taagtggctt	ttttcctaaa	360
aggggaaaaca	ggaagagana	atggcacaca	aaacaaacat	tttatattca	tatttctacc	420
tacgttaata	aaatagcatt	ttgtgaagcc	agctcaaaag	aaggcttaga	tccttttatg	480
tccatttttag	tcactaaacg	atatacnaag	tgccagaatg	caaaagggtt	gtgaacattt	540
attcaaaagc	taatataaga	tatttcacat	actcatcttt	ctg		583

<210> 204  
 <211> 589  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(589)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 204

ttttttttnt	tttttttttt	tttttttntct	ttcttttttt	ttganaatga	ggatcgagtt	60
tttcactctc	tagatagggc	atgaagaaaa	ctcatctttc	cagctttaa	ataacaatca	120
aatctcttat	gctatatcat	attttaagtt	aaactaatga	gtcactggct	tatcttctcc	180
tgaaggaaat	ctgttcattc	ttctcattca	tatagttata	tcaagtacta	ccttgcatat	240
tgagaggttt	ttcttctcta	tttacacata	tatttccatg	tgaatttgta	tcaaaccttt	300
attttcatgc	aaactagaaa	ataatgtntt	cttttgcata	agagaagaga	acaatatnag	360
cattacaaaa	ctgctcaaat	tgtttgtaa	gnntatccat	tataattagt	tnggcaggag	420
ctaatacaaa	tcacatttac	ngacnagcaa	taataaaact	gaagtaccag	ttaaatatcc	480
aaaataatta	aaggaacatt	tttagcctgg	gtataattag	ctaattcact	ttacaagcat	540
ttattnagaa	tgaattcaca	tgttattatt	ccntagccca	acacaatgg		589

&lt;210&gt; 205

&lt;211&gt; 545

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(545)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 205

tttttntttt	ttttttcagt	aataatcaga	acaatattta	tttttatatt	taaaattcat	60
agaaaagtgc	cttacattta	ataaaagttt	gtttctcaaa	gtgatcagag	gaattagata	120
tngtcttgaa	caccaatatt	aatttgagga	aaatacacca	aaatacatta	agtaaattat	180
ttaagatcat	agagcttgta	agtgaaaaga	taaaatttga	cctcagaaac	tctgagcatt	240
aaaaatccac	tattagcaaa	taaattacta	tggaacttct	gctttaattt	tgtgatgaat	300
atgggggtgc	actggtaaac	caacacattc	tgaaggatac	attacttagt	gatagattct	360
tatgtacttt	gctanatnac	gtggatatga	gttgacaagt	ttctctttct	tcaatctttt	420
aaggggcnga	ngaaatgagg	aagaaaagaa	aaggattacg	catactgttc	tttctatngg	480
aaggattaga	tatgtttcct	ttgccaatat	taaaaaata	ataatgttta	ctactagtga	540
aacc						545

&lt;210&gt; 206

&lt;211&gt; 487

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(487)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 206

tttttttttt	ttttttagtc	aagtttctna	tttttattat	aattaaagtc	ttgggtcattt	60
catttattag	ctctgcaact	tacatattta	aattaaagaa	acgttnttag	acaactgtna	120
caatttataa	atgtaagggtg	ccattattga	gtanatatat	tcctccaaga	gtggatgtgt	180
cccttctccc	accaactaat	gaancagcaa	cattagttta	attttattag	tagatnatac	240
actgctgcaa	acgctaattc	tcttctccat	ccccatgtng	atattgtgta	tatgtgtgag	300
ttggtnagaa	tgcatcanca	atctnacaat	caacagcaag	atgaagctag	gcntgggctt	360
tcggtgaaaa	tagactgtgt	ctgtctgaat	caaatagatct	gacctatcct	cggtggcaag	420
aactcttcga	accgcttcct	caaaggcngc	tgccacattt	gtggcntctn	ttgcacttgt	480



ttcaaaa

487

&lt;210&gt; 207

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(332)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 207

tgaattggct	aaaagactgc	atTTTTanaa	ctagcaactc	ttatttcttt	cctttaaaaa	60
tacatagcat	taaattccaa	atcctattta	aagacctgac	agcttgagaa	ggtcactact	120
gcatttatag	gaccttctgg	tggttctgct	gttacntttg	aantctgaca	atccttgana	180
atctttgcat	gcagaggagg	taaaaggtat	tggattttca	cagaggaana	acacagcgca	240
gaaatgaagg	ggccaggcct	actgagcttg	tccactggag	ggctcatggg	tgggacatgg	300
aaaagaaggc	agcctaggcc	ctggggagcc	ca			332

&lt;210&gt; 208

&lt;211&gt; 524

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(524)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 208

agggcggtgt	gctggaggcg	ttactgtttt	gtctcagtaa	caataaatac	aaaaagactg	60
gttgtgttcc	ggccccatcc	aaccacgaag	ttgatttctc	tttgtgtgcag	agtgactgat	120
tttaaaggac	atggagcttg	tcacaatgtc	acaatgtcac	agtgtgaagg	gcacactcac	180
tcccgcgtag	ttcacattta	gcaaccaaca	atagctcatg	agtccatact	tgtaaatact	240
tttggcagaa	tacttnttga	aacttgcaga	tgataactaa	gatccaagat	atttcccaa	300
gtaaatagaa	gtgggtcata	atattaatta	cctgttcaca	tcagcttcca	tttacaagtc	360
atgagcccag	acactgacat	caaactaagc	ccacttagac	tcctcaccac	cagtctgtcc	420
tgatcatcaga	caggaggctg	tcaccttgac	caaattctca	ccagtcaatc	atctatccaa	480
aaaccattac	ctgatccact	tccggtaatg	caccaccttg	gtga		524

&lt;210&gt; 209

&lt;211&gt; 159

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 209

gggtgaggaa	atccagagtt	gccatggaga	aaattccagt	gtcagcattc	ttgctccttg	60
tggccctctc	ctacactctg	gccagagata	ccacagtcaa	acctggagcc	aaaaaggaca	120
caaaggactc	tcgacccaaa	ctgccccaga	ccctctcca			159

&lt;210&gt; 210

&lt;211&gt; 256

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(256)  
 <223> n = A,T,C or G

<400> 210  
 actccctggc agacaaaggc agaggagaga gctctgttag ttctgtgttg ttgaactgcc 60  
 actgaatttc tttccacttg gactattaca tgccanttga gggactaatg gaaaaacgta 120  
 tggggagatt ttanccaatt tangtntgta aatgggggaga ctggggcagg cgggagagat 180  
 ttgcagggtg naaatgggan ggctggtttg ttanatgaac agggacatag gaggtaggca 240  
 ccaggatgct aatatca 256

<210> 211  
 <211> 264  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(264)  
 <223> n = A,T,C or G

<400> 211  
 acattgtttt tttgagataa agcattgaga gagctctcct taacgtgaca caatggaagg 60  
 actggaacac atacccacat ctttgttctg agggataatt ttctgataaa gtcttgctgt 120  
 atattcaagc acatatgtta tatattattc agttccatgt ttatagccta gtaaggaga 180  
 ggggagatac attcngaaaag aggactgaaa gaaatactca agtnggaaaa cagaaaaaga 240  
 aaaaaaggag caaatgagaa gcct 264

<210> 212  
 <211> 328  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(328)  
 <223> n = A,T,C or G

<400> 212  
 acccaaaaat ccaatgctga atatttggtc tcattattcc canattcttt gattgtcaaa 60  
 ggatttaatg ttgtctcagc ttgggcactt cagttaggac ctaaggatgc cagccggcag 120  
 gtttatatat gcagcaacaa tattcaagcg cgacaacagg ttattgaact tgcccggcag 180  
 ttnaatttca ttcccattga cttgggatcc ttatcatcag ccagagagat tgaaaattta 240  
 cccctacnac tctttactct ctgganaggg ccagtgggtg tagctataag cttggccaca 300  
 ttttttttct ctttattcct ttgtcaga 328

<210> 213  
 <211> 250  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature

&lt;222&gt; (1) ... (250)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 213

acttatgagc agagcgacat atccnagtgt agactgaata aaactgaatt ctctccagtt	60
taaagcattg ctactgaag ggatagaagt gactgccagg agggaaagta agccaaggct	120
cattatgcc aagganatat acatttcaat tctccaaact tcttcctcat tccaagagtt	180
ttcaatattt gcatgaacct gctgataanc catgttaana aacaaatata tctctnacct	240
tctcatcggt	250

&lt;210&gt; 214

&lt;211&gt; 444

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (444)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 214

accagaatc caatgctgaa tatttggctt cattattccc agattctttg attgtcaaag	60
gatttaagt tgtctcagct tgggcacttc agttaggacc taaggatgcc agccggcagg	120
tttatatatg cagcaacaat attcaagcgc gacaacaggt tattgaactt gcccgccagt	180
tgaatttcat tcccattgac ttgggatcct tatcatcagc canagagatt gaaaatttac	240
ccctacgact ctttactctc tggagagggc cagtgggtgg agctataagc ttggccacat	300
ttttttttcc tttattcctt tgtcagagat gcgattcatc catatgctan aaaccaacag	360
agtgaacttt acaaaattcc tataganatt gtgaataaaa ccttacctat agttgccatt	420
actttgctct ccctaataata cctc	444

&lt;210&gt; 215

&lt;211&gt; 366

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (366)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 215

acttatgagc agagcgacat atccaagtgt anactgaata aaactgaatt ctctccagtt	60
taaagcattg ctactgaag ggatagaagt gactgccagg agggaaagta agccaaggct	120
cattatgcc aagganatat acatttcaat tctccaaact tcttcctcat tccaagagtt	180
ttcaatattt gcatgaacct gctgataagc catgttgaga aacaaatata tctctgacct	240
tctcatcggt aagcagaggc tgtaggcaac atggaccata gcgaanaaaa aacttagtaa	300
tccaagctgt tttctacact gtaaccagggt ttccaaccaa ggtggaaatc tcttatactt	360
ggtgcc	366

&lt;210&gt; 216

&lt;211&gt; 260

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

<221> misc\_feature  
 <222> (1)...(260)  
 <223> n = A,T,C or G

<400> 216

ctgtataaac agaactccac tgcangaggg agggccgggc caggagaatc tccgcttgtc	60
caagacaggg gcctaaggag ggtctccaca ctgctnntaa gggctntnc atttttttat	120
taataaaaag tnnaaaaggc ctcttctcaa cttttttccc ttnggctgga aaatttaaaa	180
atcaaaaatt tcctnaagtt ntcaagctat catatatact ntatcctgaa aaagcaacat	240
aattcttctt tccctccttt	260

<210> 217  
 <211> 262  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(262)  
 <223> n = A,T,C or G

<400> 217

acctacgtgg gtaagtttan aaatgttata atttcaggaa naggaacgca tataattgta	60
tcttgctat aattttctat tttaataagg aaatagcaaa ttgggggtggg gggaatgtag	120
ggcattctac agtttgagca aaatgcaatt aaatgtggaa ggacagcact gaaaaatttt	180
atgaataatc tgtatgatta tatgtctcta gagtagattt ataattagcc acttacccta	240
atattcttca tgcttgtaaa gt	262

<210> 218  
 <211> 205  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(205)  
 <223> n = A,T,C or G

<400> 218

accaaggtgg tgcattaccg gaantggatc aangacacca tcgtggccaa cccctgagca	60
cccctatcaa ctcccttttg tagtaaaactt ggaaccttgg aaatgaccag gccaaagactc	120
aggcctcccc agttctactg accttgtcc ttangntna ngtccagggt tgctagggaa	180
anaaatcagc agacacaggt gtaa	205

<210> 219  
 <211> 114  
 <212> DNA  
 <213> Homo sapien

<400> 219

tactgttttg tctcagtaac aataaatata aaaagactgg ttgtgttccg gcccatcca	60
accacgaagt tgatttctct tgtgtgcaga gtgactgatt ttaaaggaca tgga	114

<210> 220  
 <211> 93

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 220

actagccagc acaaaaggca gggtagcctg aattgctttc tgctctttac atttctttta 60  
 aaataagcat ttagtgctca gtcctactg agt 93

&lt;210&gt; 221

&lt;211&gt; 167

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (167)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 221

actangtgca ggtgcgca aatatttgct gatattccct tcattcttga ttccatgagg 60  
 tcttttgccc agcctgtggc tctactgtag taagtttctg ctgatgagga gccagnatgc 120  
 cccccactac cttccctgac gtccccaana aatcacccaa cctctgt 167

&lt;210&gt; 222

&lt;211&gt; 351

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 222

agggcggtgt gcgagggtgt gtactgacct cattagtagg aggatgcatt ctggcaccac 60  
 gttcttcacc tgtcccccaa tcttataaag gccatactgc ataaagtcaa caacagataa 120  
 atgtttgctg aattaaagga tggatgaaaa aaattaataa tgaatttttg cataatccaa 180  
 ttttctcttt tatatttcta gaagaagttt ctttgagcct attagatccc gggaatcttt 240  
 taggtgagca tgattagaga gcttgtaggt tgcttttaca tatatctggc atatttgagt 300  
 ctcgtatcaa aacaatagat tggtaaaggt ggtattattg tattgataag t 351

&lt;210&gt; 223

&lt;211&gt; 383

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (383)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 223

aaaacaaaca aacaaaaaaa acaattcttc attcagaaaa attatcttag ggactgatat 60  
 tggtaattat ggtcaattta atwrtrttkt ggggcatttc cttacattgt cttgacaaga 120  
 ttaaaatgtc tgtgccaaaa ttttgatatt tatttgagga cttcttatca aaagtaatgc 180  
 tgccaaagga agtctaagga attagtagtg ttcccmcac ttgtttggag tgtgctattc 240  
 taaaagattt tgatttcttg gaatgacaat tatattttta ctttggtggg ggaaanagtt 300  
 ataggaccac agtcttcact tctgatactt gtaaattaat cttttattgc acttgttttg 360  
 accattaagc tatatgttta aaa 383

&lt;210&gt; 224

&lt;211&gt; 320

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 224

cccctgaagg	cttcttggtta	gaaaatagta	cagttacaac	caataggaac	aacaaaaaga	60
aaaagtttgt	gacattgtag	tagggagtgt	gtacccctta	ctcccatca	aaaaaaaaat	120
ggatacatgg	ttaaaggata	raagggcaat	attttatcat	atgttctaaa	agagaaggaa	180
gagaaaatac	tactttctcr	aaatggaagc	ccttaaaggt	gctttgatac	tgaaggacac	240
aaatgtggcc	gtccatcctc	ctttaragtt	gcatgacttg	gacacggtaa	ctggtgcagt	300
tttaractcm	gcattgtgac					320

&lt;210&gt; 225

&lt;211&gt; 1214

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 225

gaggactgca	gccccgactc	gcagccctgg	caggcggcac	tgggtcatgga	aaacgaattg	60
ttctgctcgg	gcgtcctggt	gcattccgag	tgggtgctgt	cagccgcaca	ctgtttccag	120
aactcctaca	ccatcgggct	gggcctgcac	agtcttgagg	ccgaccaaga	gccagggagc	180
cagatggtgg	aggccagcct	ctccgtacgg	caccagagt	acaacagacc	cttgctcgct	240
aacgacctca	tgctcatcaa	gttgacgaa	tccgtgtccg	agtctgacac	catccggagc	300
atcagcattg	cttcgcagtg	ccctaccgcg	gggaactctt	gcctcgtttc	tggctggggg	360
ctgctggcga	acggcagaat	gcctaccgtg	ctgcagtgcg	tgaacgtgtc	ggtggtgtct	420
gaggaggtct	gcagtaagct	ctatgacccg	ctgtaccacc	ccagcatggt	ctgcgccggc	480
ggagggcaag	accagaagga	ctcctgcaac	ggtgactctg	gggggcccc	gatctgcaac	540
gggtacttgc	agggccttgt	gtctttcgga	aaagccccgt	gtggccaagt	tggcgtgcca	600
ggtgtctaca	ccaacctctg	caaattcact	gagtggatag	agaaaaccgt	ccaggccagt	660
taactctggg	gactgggaac	ccatgaaatt	gacccccaaa	tacatcctgc	ggaaggaatt	720
caggaatatc	tgttcccagc	ccctcctccc	tcaggcccag	gagtcagacc	ccccagcccc	780
tcctccctca	aaccaagggt	acagatcccc	agccccctct	ccctcagacc	caggagtcca	840
gacccccag	cccctcctcc	ctcagaccca	ggagtccagc	ccctcctccc	tcagaccag	900
gagtcagac	ccccagccc	ctcctcctcc	agacccaggg	gtccaggccc	ccaacccctc	960
ctccctcaga	ctcagaggtc	caagccccca	acccctcctt	ccccagaccc	agaggtccag	1020
gtcccagccc	ctcctcctcc	agacccagcg	gtccaatgcc	acctagactc	tcctgttaca	1080
cagtgccccc	ttgtggcacg	ttgacccaac	cttaccagtt	ggtttttcat	tttttgtccc	1140
tttcccctag	atccgaaat	aaagtctaag	agaagcgcaa	aaaaaaaaaa	aaaaaaaaaa	1200
aaaaaaaaaa	aaaa					1214

&lt;210&gt; 226

&lt;211&gt; 119

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 226

accagtatg	tgcagggaga	cggaacccca	tgtgacagcc	cactccacca	gggttcccaa	60
agaacctggc	ccagtcataa	tcattcatcc	tgacagtggc	aataatcacg	ataaccagt	119

&lt;210&gt; 227

&lt;211&gt; 818

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 227

acaattcata	gggacgacca	atgaggacag	ggaatgaacc	cggtctctccc	ccagccctga	60
tttttgctac	atatggggtc	ccttttcatt	ctttgcaaaa	acactgggtt	ttctgagaac	120
acggacgggt	cttagcacia	tttgtgaaat	ctgtgtaraa	ccgggctttg	caggggagat	180
aattttcctc	ctctggagga	aagggtggtga	ttgacaggca	gggagacagt	gacaaggcta	240
gagaaagcca	cgctcggcct	tctctgaacc	aggatggaac	ggcagacccc	tgaaaacgaa	300
gcttgctccc	ttccaatcag	ccacttctga	gaacccccat	ctaacttcct	actggaaaag	360
agggcctcct	caggagcagt	ccaagagttt	tcaaagataa	cgtagacaact	accatctaga	420
ggaaagggtg	caecctcagc	agagaagccg	agagcttaac	tctggtcggt	tccagagaca	480
acctgctggc	tgtcttggga	tgcgcccagc	ctttgagagg	ccactacccc	atgaacttct	540
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aaagccattc	ccacaaatcc	agaccatacc	atgaagcaac	gagacccaaa	cagtttggct	720
caagaggata	tgaggactgt	ctcagcctgg	ctttgggctg	acaccatgca	cacacacaag	780
gtccacttct	aggttttccag	cctagatggg	agtcgtgt			818

&lt;210&gt; 228

&lt;211&gt; 744

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 228

actggagaca	ctgttgaact	tgatcaagac	ccagaccacc	ccaggtctcc	ttcgtgggat	60
gtcatgacgt	ttgacatacc	tttggaaacga	gcctcctcct	tggaaagatgg	aagaccgtgt	120
tcgtggccga	cctggcctct	cctggcctgt	ttcttaagat	gcggagtcac	atttcaatgg	180
taggaaaagt	ggcttcgtaa	aatagaagag	cagtcactgt	ggaactacca	aatggcgaga	240
tgctcggtag	acattggggg	gctttgggat	aaaagattta	tgagccaact	attctctggc	300
accagattct	aggccagttt	gttccactga	agcttttccc	acagcagtc	acctctgcag	360
gctggcagct	gaatggcctt	ccgggtggct	tgtggcaaga	tcacactgag	atcgatgggt	420
gagaaggcta	ggatgcttgt	ctagtgttct	tagctgtcac	gttggctcct	tccaggttgg	480
ccagacgggt	ttggccactc	ccttctaata	cacaggcgcc	ctcctgggtga	cagtgaaccg	540
ccgtgggtat	ccttggccca	ttccagcagt	cccagttatg	catttcaagt	ttggggtttg	600
ttcttttctg	taatgttctt	ctgtgttctc	agctgtcttc	atttctctgg	ctaagcagca	660
ttgggagatg	tggaccagag	atccactcct	taagaaccag	tggcgaaaga	cactttcttt	720
cttcaactct	aagtagctgg	tggt				744

&lt;210&gt; 229

&lt;211&gt; 300

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 229

cgagtctggg	ttttgtctat	aaagtttcat	ccctcctttt	ctcatccaaa	tcatgtgaac	60
cattacacat	cgaaataaaa	gaaaggtggc	agacttgccc	aacgccaggc	tgacatgtgc	120
tgcagggttg	ttgtttttta	attattattg	ttagaaacgt	cacccacagt	ccctgttaat	180
ttgtatgtga	cagccaactc	tgagaaggct	ctatttttcc	acctgcagag	gatccagctc	240
cactaggctc	ctccttgccc	tcacactgga	gtctccgcca	gtgtgggtgc	ccactgacat	300

&lt;210&gt; 230

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 230

cagcagaaca	aatacaata	tgaagagtgc	aaagatctca	taaaatctat	gctgaggaat	60
gagcgacagt	tcaaggagga	gaagcttgca	gagcagctca	agcaagctga	ggagctcagg	120

caatataaag tcctggttca cactcaggaa cgagagctga cccagttaag ggagaagttg 180  
 cgggaagggga gagatgcctc cctctcattg aatgagcatc tccaggccct cctcactccg 240  
 gatgaaccgg acaagtccca ggggcaggac ctccaagaaa cagacctcgg ccgcgaccac 300  
 g 301

<210> 231  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 231  
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 caggaactcc aagtccacat ccttggaac tggggacttg cgcagggttag ccttgaggat 120  
 ggcaacacgg gactttctcat caggaagtgg gatgtagatg agctgatcaa gacggccagg 180  
 tctgaggatg gcaggatcaa tgatgtcagg ccggttggtta ccgccaatga tgaacacatt 240  
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 c 301

<210> 232  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 232  
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 ggcgacagcg gggcttcctg attctggaat ataactttgt gtaaattaac agccacctat 120  
 agaagagtcc atctgctgtg aaggagagac agagaactct gggttccgct gcctgttcca 180  
 cgtgctgtac caagtgtctg tgccagcctg ttacctgttc tctactgaaa tctggctaata 240  
 gctcttctgt atcacttctg attctgacaa tcaatcaatc aatggcctag agcactgact 300  
 g 301

<210> 233  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 233  
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 cctagaagtt acagagcatc tagctggtgc gctggcacc cttggcctcac acagactccc 180  
 gagtagctgg gactacaggc acacagtcac tgaagcaggc cctgtttagca attctatgag 240  
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 c 301

<210> 234  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 234  
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 tcaatttcag caacatactt ctcaaattct tcaggattta aaatcttgag ggattgatct 180  
 cgcctcatga cagcaagttc aatgtttttg ccacctgact gaaccacttc caggagtgcc 240  
 ttgatcacca gcttaatggt cagatcatct gcttcaatgg cttcgtcagt atagttcttc 300



t

301

&lt;210&gt; 235

&lt;211&gt; 283

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 235

tggggctgtg catcaggcgg gtttgagaaa tattcaattc tcagcagaag ccagaatttg	60
aattccctca tcttttaggg aatcatttac caggtttgga gaggattcag acagctcagg	120
tgctttcact aatgtctctg aacttctgtc cctctttgtt catggatagt ccaataaata	180
atgttatctt tgaactgatg ctcataggag agaataaag aactctgagt gatatcaaca	240
ttagggattc aaagaaatat tagattttaag ctcacactgg tca	283

&lt;210&gt; 236

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 236

aggtcctcca ccaactgcct gaagcacggg taaaattggg aagaagtata gtgcagcata	60
aatactttta atcgcacag atttccctaa cccacatgca atcttcttca ccagaagagg	120
tcggagcagc atcattaata ccaagcagaa tgcgtaatag ataaatacaa tggatatag	180
tgggtagacg gcttcatgag tacagtgtac tgtggtatcg taatctggac ttgggttgta	240
aagcatcgtg taccagtcag aaagcatcaa tactcgacat gaacgaatat aaagaacacc	300
a	301

&lt;210&gt; 237

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 237

cagtggtagt ggtgggtggac gtggcggttg tcgtgggtgcc ttttttggtg cccgtcacaa	60
actcaatttt tgttcgctcc tttttggcct ttccaattt gtccatctca attttctggg	120
ccttggttaa tgcctcatag taggagtcct cagaccagcc atggggatca aacatatcct	180
ttgggtagtt ggtgccaaagc tcgtcaatgg cacagaatgg atcagcttct cgtaaactta	240
gggttccgaa attctttctt cctttggata atgtagtcca tatccattcc ctcttttatt	300
t	301

&lt;210&gt; 238

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 238

gggcagggtt tttttttttt ttttttgatg gtgcagaccc ttgctttatt tgtctgactt	60
gttcacagtt cagccccctg ctcagaaaac caacgggcca gctaaggaga ggaggaggca	120
ccttgagact tccggagtcg aggtctctcca ggggtccccca gccatcaat cattttctgc	180
acccccctgc tgggaagcag ctccctgggg ggtgggaatg ggtgactaga agggatttca	240
gtgtgggacc cagggtctgt tcttcacagt aggaggtgga agggatgact aatttcttta	300
t	301

&lt;210&gt; 239

&lt;211&gt; 239

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 239

ataagcagct aggggaattct ttatttagta atgtcctaac ataaaagtgc acataactgc	60
ttctgtcaaa ccatgatact gagctttgtg acaaccaga aataactaag agaaggcaaa	120
cataatacct tagagatcaa gaaacattta cacagttcaa ctgtttaaaa atagctcaac	180
attcagccag tgagtagagt gtgaatgcc aacatacag tatacaggte cttcaggga	239

&lt;210&gt; 240

&lt;211&gt; 300

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 240

ggtcctaattg aagcagcagc ttccacattt taacgcaggt ttacgggtgat actgtccttt	60
gggatctgcc ctccagtggg accttttaag gaagaagtgg gcccaagcta agttccacat	120
gctgggtgag ccagatgact tctgttccct ggtcactttc ttcaatgggg cgaatggggg	180
ctgccaggtt tttaaaatca tgcttcatct tgaagcacac ggtcacttca ccctcctcac	240
gctgtgggtg taatttgatg aaaataccca ctttgttggc ctttctgaag ctataatgtc	300

&lt;210&gt; 241

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 241

gaggtctggt gctgaggtct ctgggctagg aagaggagtt ctgtggagct ggaagccaga	60
cctcttttga ggaactcca gcagctatgt tgggtgtctt gaggggaatgc aacaaggctg	120
ctcctccatg tattggaaaa ctgcaaactg gactcaactg gaaggaagtg ctgctgccag	180
tgtgaagaac cagcctgagg tgacagaaac ggaagcaaac aggaacagcc agtcttttct	240
tcctcctcct gtcatacggg ctctctcaag catcctttgt tgtcaggggc ctaaaaggga	300

&lt;210&gt; 242

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 242

ccgaggtcct gggatgcaac caatcactct gtttcacgtg acttttatca ccatacaatt	60
tgtggcattt cctcattttc tacattgtag aatcaagagt gtaataaat gtatatcgat	120
gtcttcaaga atatatcatt cttttttcac tagaaccat tcaaaatata agtcaagaat	180
cttaatatca acaaatatat caagcaaact ggaaggcaga ataactacca taatttagta	240
taagtaccca aagttttata aatcaaaagc cctaatagata accattttta gaattcaatc	300

&lt;210&gt; 243

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 243

aggtaagtcc cagtttgaag ctcaaaagat ctggtatgag cataggctca tcgacgacat	60
ggtggcccaa gctatgaaat cagagggagg cttcatctgg gcctgtaaaa actatgatgg	120

tgacgtgcag tcggactctg tggcccaagg gtatggctct ctccggcatga tgaccagcgt 180  
 gctggtttgt ccagatggca agacagtaga agcagaggct gcccacggga ctgtaaccgc 240  
 tcactaccgc atgttccaga aaggacagga gacgtccacc aatcccattg cttccatttt 300  
 t 301

&lt;210&gt; 244

&lt;211&gt; 300

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 244

gctggtttgc aagaatgaaa tgaatgattc tacagctagg acttaacctt gaaatggaaa 60  
 gtcacgcaat cccatttgcg ggatctgtct gtgcacatgc ctctgtagag agcagcattc 120  
 ccagggacct tggaaacagt tgacactgta aggtgcttgc tccccaagac acatcctaaa 180  
 aggtgttgta atggtgaaaa cgtcttctct ctttattgcc ccttcttatt tatgtgaaca 240  
 actgtttgtc ttttgtgtat cttttttaaa ctgtaaagtt caattgtgaa aatgaatatt 300

&lt;210&gt; 245

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 245

gtctgagtat ttaaaatggt attgaaatta tccccaacca atgttagaaa agaaagaggt 60  
 tatatactta gataaaaaat gaggtgaatt actatccatt gaaatcatgc tcttagaatt 120  
 aaggccagga gatattgtca ttaatgtara ctccaggaca cttagagtata gcagccctat 180  
 gttttcaaag agcagagatg caattaaata ttgttttagca tcaaaaaggc cactcaatac 240  
 agctaataaa atgaaagacc taatttctaa agcaattctt tataatttac aaagttttaa 300  
 g 301

&lt;210&gt; 246

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 246

ggctctgtct acaatgcctg cttcttgaaa gaagtcggca ctttctagaa tagctaaata 60  
 acctgggctt attttaaaga actatttgta gctcagattg gttttcctat ggctaaaata 120  
 agtgcttctt gtgaaaatta aataaaacag ttaattcaaa gccttgatat atgttaccac 180  
 taacaatcat actaaatata ttttgaagta caaagtttga catgctctaa agtgacaacc 240  
 caaatgtgtc ttacaaaaca cgttcttaac aaggtatgct ttacactacc aatgcagaaa 300  
 c 301

&lt;210&gt; 247

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 247

aggtcctttg gcagggtcga tggatcagag ctcaaactgg agggaaaggc atttcgggta 60  
 gcctaagagg gcgactggcg gcagcacaac caaggaaggc aagggtgttt cccccacgct 120  
 gtgtcctgtg ttcagggtgcg acacacaatc ctcatgggaa caggatcacc catgcgctgc 180  
 ccttgatgat caaggttggg gcttaagtgg attaaggagg gcaagttctg gggtccttgc 240  
 cttttcaaac catgaagtca ggctctgtat ccctcctttt cctaactgat attctaacta 300  
 a 301

<210> 248  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 248

aggtccttgg agatgccatt tcagccgaag gactcttctw ttcggaagta caccctcact	60
attaggaaga ttcttagggg taatttttct gaggaaggag aactagccaa cttaagaatt	120
acaggaagaa agtgggttgg aagacagcca aagaaataaa agcagattaa attgtatcag	180
gtacattcca gcctgttggc aactccataa aaacatttca gattttaatc ccgaatttag	240
ctaattgagac tggatttttg ttttttatgt tgtgtgtcgc agagctaaaa actcagttcc	300
c	301

<210> 249  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 249

gtccagagga agcacctggt gctgaactag gcttgccctg ctgtgaactt gcacttggag	60
ccctgacgct gctgttctcc ccgaaaaacc cgaccgacct ccgcgatctc cgccccgcc	120
ccagggagac acagcagtga ctacagagctg gtgcacact gtgcctccct cctcaccgcc	180
catcgtaatg aattattttg aaaattaatt ccaccatcct ttcagattct ggatggaaag	240
actgaatctt tgactcagaa ttgtttgctg aaaagaatga tgtgactttc ttagtcattt	300
a	301

<210> 250  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 250

ggtctgtgac aaggacttgc aggctgtggg aggcaagtga cccttaacac tacacttctc	60
cttatcttta ttggcttgat aaacataatt atttctaaca ctagcttatt tccagttgcc	120
cataagcaca tcagtacttt tctctggctg gaatagtaaa ctaaagtatg gtacatctac	180
ctaaaagact actatgtgga ataatacata ctaatgaagt attacatgat ttaaagacta	240
caataaaacc aaacatgctt ataacattaa gaaaaacaat aaagatacat gattgaaacc	300
a	301

<210> 251  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 251

gccgaggtcc tacatttggc ccagtttccc cctgcatect ctccagggcc cctgcctcat	60
agacaacctc atagagcata ggagaactgg ttgccctggg ggcaggggga ctgtctggat	120
ggcaggggtc ctcaaaaatg ccactgtcac tgccaggaaa tgcttctgag cagtacacct	180
cattgggatc aatgaaaagc ttcaagaaat cttcaggctc actctcttga aggcccgga	240
cctctggagg ggggcagtgg aatcccagct ccaggacgga tctgtcgaag aagatatact	300
c	301

<210> 252  
 <211> 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 252

```

gcaaccaatc actctgtttc acgtgacttt tatcaccata caatttgtgg catttcctca      60
ttttctacat tgtagaatca agagtgtaaa taaatgtata tcgatgtctt caagaatata      120
tcattccttt ttcactagga acccattcaa aatataagtc aagaatctta atatcaacaa      180
atatatcaag caaactggaa ggcagaataa ctaccataat ttagtataag tacccaaagt      240
tttataaatc aaaagcccta atgataacca tttttagaat tcaatcatca ctgtagaatc      300
a

```

&lt;210&gt; 253

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 253

```

ttccctaaga agatgttatt ttgttggggt ttgttcccc tccatctcga ttctcgtacc      60
caactaaaaa aaaaaataa agaaaaaatg tgctgcgttc tgaaaaataa ctcccttagct      120
tggtctgatt gttttcagac cttaaaatat aaacttggtt cacaagcttt aatccatgtg      180
gatttttttt cttagagaac cacaaaacat aaaaggagca agtcggactg aatacctgtt      240
tccatagtgc ccacagggtt ttcctcacat tttctccata ggaaaatgct ttttcccaag      300
g

```

&lt;210&gt; 254

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 254

```

cgctgcgcct ttcccttggg ggagggggcaa ggccagaggg ggtccaagtg cagcacgagg      60
aacttgacca attcccttga agcgggtggg ttaaaccctg taaatgggaa caaaatcccc      120
ccaaatctct tcattctacc ctgggtggact cctgactgta gaattttttg gttgaaacaa      180
gaaaaaataa agcttttggg cttttcaagg ttgcttaaca ggtactgaaa gactggcctc      240
acttaactg agccaggaaa agctgcagat ttattaatgg gtgtgttagt gtgcagtgcc      300
t

```

&lt;210&gt; 255

&lt;211&gt; 302

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 255

```

agcttttttt tttttttttt tttttttttt ttcattaaaa aatagtgtc tttattataa      60
attactgaaa tgtttctttt ctgaatataa atataaatat gtgcaaagt tgaactggat      120
tgggattttt ttgagttctt caagcatctc ctaataccct caagggcctg agtagggggg      180
aggaaaaagg actggaggtg gaatctttat aaaaaacaag agtgattgag gcagattgta      240
aacattatta aaaaacaaga aacaaacaaa aaaatagaga aaaaaaccac cccaacacac      300
aa

```

&lt;210&gt; 256

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 256

gttccagaaa	acattgaagg	tggcttccca	aagtctaact	agggatacce	cctctagcct	60
aggaccctcc	tccccacacc	tcaatccacc	aaaccatcca	taatgcaccc	agataggccc	120
acccccaaaa	gcctggacac	cttgagcaca	cagttatgac	caggacagac	tcatctctat	180
aggcaaatag	ctgctggcaa	actggcatta	cctggtttgt	ggggatgggg	gggcaagtgt	240
gtggcctctc	ggcctggtta	gcaagaacat	tcagggtagg	cctaagttan	tcgtgttagt	300
t						301

&lt;210&gt; 257

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 257

gttgtggagg	aactctggct	tgctcattaa	gtcctactga	ttttcactat	cccctgaatt	60
tccccactta	tttttgtctt	tcactatcgc	aggccttaga	agaggtctac	ctgcctccag	120
tcttacctag	tccagtctac	cccctggagt	tagaatggcc	atcctgaagt	gaaaagtaat	180
gtcacattac	tcccttcagt	gatttcttgt	agaagtgcc	atccctgaat	gccaccaaga	240
tcttaattct	cacatcttta	atcttatctc	tttgactcct	ctttacaccg	gagaaggctc	300
c						301

&lt;210&gt; 258

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 258

cagcagtagt	agatgccgta	tgccagcacg	cccagcactc	ccaggatcag	caccagcacc	60
agggggcccag	ccaccaggcg	cagaagcaag	ataaacagta	ggctcaagac	cagagccacc	120
cccaggggcaa	caagaatcca	ataccaggac	tgggcaaaat	cttcaaagat	cttaacactg	180
atgtctcggg	cattgaggct	gtcaataana	cgctgatccc	ctgctgtatg	gtggtgtcat	240
tggtgatccc	tgggagcgcc	ggtggagtaa	cgttgggtcca	tggaaagcag	cgcccacaac	300
t						301

&lt;210&gt; 259

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 259

```

tcatatatgc aaacaaatgc agactangcc tcaggcagag actaaaggac atctcttggg      60
gtgtcctgaa gtgatttgga cccctgaggg cagacaccta agtaggaatc ccagtgggaa      120
gcaaagccat aaggaagccc aggattcctt gtgatcagga agtgggcccag gaaggtctgt      180
tccagctcac atctcatctg catgcagcac ggaccggatg cgeccactgg gtcttggctt      240
ccctcccatc ttctcaagca gtgtccttgt tgagccattt gcateccttg ctccagggtg      300
c

```

```

<210> 260
<211> 301
<212> DNA
<213> Homo sapien

```

```

<400> 260
ttttttttct ccctaaggaa aaagaaggaa caagtctcat aaaaccaa at aagcaatggt      60
aaggtgtctt aacttgaaaa agattaggag tcaactggtt acaagttata attgaatgaa      120
agaactgtaa cagccacagt tggccatttc atgccaatgg cagcaaacaa caggattaac      180
tagggcaaaa taaataagtg tgtggaagcc ctgataagtg cttaataaac agactgattc      240
actgagacat cagtacctgc cggggcggcc gtcgagccg aattctgcag atatccatca      300
c

```

```

<210> 261
<211> 301
<212> DNA
<213> Homo sapien

```

```

<400> 261
aaatatctga gcaaattcctg taactaatgt gtctccataa aaggctttga actcagtgaa      60
tctgcttcca tccacgattc tagcaatgac ctctcggaca tcaaagctcc tcttaaggtt      120
agcaccaact attccatata attcatcagc aggaaataaa ggctcttcag aaggttcaat      180
ggtgacatcc aattttcttct gataatttag attcctcaca accttcctag ttaagtgaag      240
ggcatgatga tcatccaaag cccagtgggc acttactcca gactttctgc aatgaagatc      300
a

```

```

<210> 262
<211> 301
<212> DNA
<213> Homo sapien

```

```

<400> 262
gaggagagcc tggtacagca tttgtaagca cagaatactc caggagtatt tgtaattgtc      60
tgtgagcttc ttgccgcaag tctctcagaa atttaaaaag atgcaaatec ctgagtcacc      120
cctagacttc ctaaaccaga tctctggggg ctggaacctg gcactctgca tttgtaatga      180
gggctttctg gtgcacacct aattttgtgc atctttgccc taaatcctgg attagtcccc      240
catcattacc cccacattat aatgggatag attcagagca gatactctcc agcaaagaat      300
c

```

```

<210> 263
<211> 301
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(301)
<223> n = A,T,C or G

```

&lt;400&gt; 263

tttagcttgt	ggtaaatgac	tcacaaaact	gatttttaaaa	tcaagttaat	gtgaattttg	60
aaaattacta	cttaatccta	attcacaata	acaatggcat	taaggtttga	cttgagttgg	120
ttcttagtat	tatttatggt	aaataggctc	ttaccacttg	caaataactg	gccacatcat	180
taatgactga	cttcccagta	aggctctcta	aggggtaagt	angaggatcc	acaggatttg	240
agatgctaag	gccccagaga	tcgtttgatc	caaccctctt	attttcagag	gggaaaatgg	300
g						301

&lt;210&gt; 264

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 264

aaagacgtta	aaccactcta	ctaccacttg	tggaactctc	aaagggtaaa	tgacaaascc	60
aatgaatgac	tctaaaaaca	atattttacat	ttaatggttt	gtagacaata	aaaaaacaag	120
gtggatagat	ctagaattgt	aacattttta	gaaaaccata	scatttgaca	gatgagaaag	180
ctcaattata	gatgcaaagt	tataactaaa	ctactatagt	agtaaagaaa	tacatttcac	240
acccttcata	taaattcact	atcttggctt	gaggcactcc	ataaaatgta	tcacgtgcat	300
a						301

&lt;210&gt; 265

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 265

tgcccaagtt	atgtgtaagt	gtatccgcac	ccagaggtaa	aactacactg	tcctctttgt	60
cttcttgtga	cgcagtat	cttctctggg	gagaagccgg	gaagtcttct	cctggctcta	120
catattcttg	gaagtctcta	atcaactttt	gttccatttg	tttcatttct	tcaggaggga	180
ttttcagttt	gtcaacatgt	tctctaacaa	cacttgccca	tttctgtaaa	gaatccaaag	240
cagtccaagg	ctttgacatg	tcaacaacca	gcataactag	agtatccttc	agagatacgg	300
c						301

&lt;210&gt; 266

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 266

taccgtctgc	ccttcctccc	atccaggcca	tctgcgaate	tacatgggtc	ctcctattcg	60
acaccagatc	actctttcct	ctaccacag	gcttgctatg	agcaagagac	acaacctcct	120
ctcttctgtg	ttccagcttc	tttctctgtt	cttcccaccc	cttaagttct	attcctgggg	180
atagagacac	caataccat	aacctctctc	ctaagcctcc	ttataacca	gggtgcacag	240
cacagactcc	tgacaactgg	taaggccaat	gaactgggag	ctcacagctg	gctgtgcctg	300
a						301

&lt;210&gt; 267

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 267

aaagagcaca	ggccagctca	gcctgccctg	gccatctaga	ctcagcctgg	ctccatgggg	60
------------	------------	------------	------------	------------	------------	----



```

gttctcagtg ctgagtcctat ccaggaaaag ctcacctaga ccttctgagg ctgaatcttc      120
atcctcacag gcagcttctg agagcctgat attcctagcc ttgatggctt ggagtaaagc      180
ctcattctga ttctctctct tcttttcttt caagttggct ttctcacat ccctctgttc      240
aatcgcttc agcttgtctg ctttagccct catttcaga agcttcttct ctttggcatc      300
t                                                                                   301

```

&lt;210&gt; 268

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 268

```

aatgtctcac tcaactactt cccagcctac cgtggcctaa ttctgggagt tttcttctta      60
gatcttggga gagctgggtc ttctaaggag aaggaggaag gacagatgta accttggatc      120
tcgaagagga agtctaattg aagtaattag tcaacgggtc ttgtttagac tcttgggaata      180
tgctgggtgg ctgagtgagc ccttttggag aaagcaagta ttattcttaa ggagtaacca      240
cttcccattg ttctactttc taccatcatc aattgtatat tatgtattct ttggagaact      300
a                                                                                   301

```

&lt;210&gt; 269

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 269

```

taacaatata cactagctat ctttttaact gtccatcatt agcaccaatg aagattcaat      60
aaaattacct ttattcacac atctcaaaac aattctgcaa attcttagtg aagtttaact      120
atagtcacag acctaaata ttcacattgt tttctatgtc tactgaaaat aagttcacta      180
cttttctgga tattctttac aaaatcttat taaaattcct ggtattatca cccccaatta      240
tacagtagca caaccacctt atgtagtttt tacatgatag ctctgtagaa gtttcacatc      300
t                                                                                   301

```

&lt;210&gt; 270

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 270

```

cattgaagag cttttgcgaa acatcagaac acaagtgtt ataaaattaa ttaagcctta      60
cacaagaata catattcctt ttatttctaa ggagttaaac atagatgtag ctgatgtgga      120
gagcttgctg gtgcagtgca tattggataa cactattcat ggccgaattg atcaagtcaa      180
ccaactcctt gaactggatc atcagaagaa ggggtggtgca cgatatactg cactagataa      240
tggaccaacc aactaaattc tctcaccagg ctgtatcagt aaactggctt aacagaaaac      300
a                                                                                   301

```

&lt;210&gt; 271

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 271

```

aaaaggttct cataagatta acaatttaaa taaatatttg atagaacatt ctttctcatt      60
tttatagctc atcttttaggg ttgatattca gttcatgctt cccttgctgt tcttgatcca      120
gaattgcaat cacttcatca gcctgtattc gctccaattc tctataaagt ggggtccaagg      180
tgaaccacag agccacagca cacctctttc ccttggtgac tgccttcacc ccatganggt      240
tctctcctcc agatganaac tgatcatgcg cccacatttt gggttttata gaagcagtca      300
c

```

&lt;210&gt; 272

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 272

```

taaattgcta agccacagat aacaccaatc aaatggaaca aatcactgtc ttcaaagtgc      60
ttatcagaaa accaaatgag cctggaatct tcataatacc taaacatgcc gtatttagga      120
tccaataatt ccctcatgat gagcaagaaa aattctttgc gcacccctcc tgcattccaca      180
gcatcttctc aaacaaatat aaccttgagt ggcttcttgt aatctatggt ctttggtttc      240
ctaaggactt ccattgcatc tctacaata ttttctctac gcaccactag aattaagcag      300
g

```

&lt;210&gt; 273

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 273

```

acatgtgtgt atgtgtatct ttgggaaaaa aanaagacat cttgtttayt atttttttgg      60
agagangctg ggacatggat aatcacwtaa tttgctayta tyactttaat ctgactygaa      120
gaaccgtcta aaaataaaat ttaccatgct dtatattcct tatagtatgc ttatttcacc      180
ttytttctgt ccagagagag tatcagtgc ananatttma ggggtgaamac atgmattgggt      240
gggacttnty tttaacngagm accctgcccg sgcgcctctg makcngantt ccgcsananc      300
t

```

&lt;210&gt; 274

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 274

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cttatatact ctttctcaga ggcaaaagag gagatgggta atgtagacaa ttctttgagg      60
aacagtaaat gattattaga gagaangaat ggaccaagga gacagaaatt aacttgtaaa      120
tgattctctt tggaatctga atgagatcaa gaggccagct ttagcttggt gaaaagtcca      180
tctaggtatg gttgcattct cgtcttcttt tctgcagtag ataatgaggt aaccgaaggc      240
aattgtgctt cttttgataa gaagctttct tggatcatatc aggaaattcc aganaaaagtc      300

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c

301

<210> 275  
 <211> 301  
 <212> DNA  
 <213> Homo sapien  
  
 <220>  
 <221> misc\_feature  
 <222> (1)...(301)  
 <223> n = A,T,C or G

<400> 275  
 tcggtgtcag cagcacgtgg cattgaacat tgcaatgtgg agcccaaacc acagaaaatg 60  
 ggggtgaaatt ggccaacttt ctattaactt atgttgccaa ttttgccacc aacagtaagc 120  
 tggcccttct aataaaagaa aattgaaagg tttctcacta aacggaatta agtagtggag 180  
 tcaagagact cccaggcctc agcgtacctg cccggggcggc cgctcgaagc cgaattctgc 240  
 agatatccat cacactggcg gncgctcgan catgcatcta gaaggnccaa ttcgccttat 300  
 a 301

<210> 276  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 276  
 tgtacacata ctcaataaat aaatgactgc attgtggtat tattactata ctgattatat 60  
 ttatcatgtg acttctaatt agaaaatgta tccaaaagca aaacagcaga tatacaaaat 120  
 taaagagaca gaagatagac attaacagat aaggcaactt atacattgag aatccaaatc 180  
 caatacatTT aaacatttgg gaaatgaggg ggacaaatgg aagccagatc aaatttgtgt 240  
 aaaactattc agtatgtttc ccttgcttca tgtctgagaa ggctctcctt caatggggat 300  
 g 301

<210> 277  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(301)  
 <223> n = A,T,C or G

<400> 277  
 tttgttgatg tcagtatttt attacttgcg ttatgagtgc tcacctggga aattctaaag 60  
 atacagagga cttggaggaa gcagagcaac tgaatttaat ttaaaagaag gaaaacattg 120  
 gaatcatggc actcctgata ctttcccaaa tcaacactct caatgccccca cctcgtcct 180  
 caccatagtg gggagactaa agtggccacg gatttgccct anggtgtgcag tgcgttctga 240  
 gttcncgtgc gattacatct gaccagtctc ctttttccga agtccttccg ttcaatcttg 300  
 c 301

<210> 278  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 278

taccactaca ctccagcctg ggcaacagag caagacctgt ctcaaagcat aaaatggaat	60
aacatatcaa atgaaacagg gaaaatgaag ctgacaattt atggaagcca gggcttgctca	120
cagtctctac tggttattatg cattacctgg gaatttatat aagcccttaa taataatgcc	180
aatgaacatc tcatgtgtgc tcacaatgtt ctggcactat tataagtgtc tcacagggtt	240
tatgtgttct tcgtaacttt atggantagg tactcggcgc cgaacacgct aagccgaatt	300
c	301

&lt;210&gt; 279

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 279

aaagcaggaa tgacaaagct tgcttttctg gtatgttcta ggtgtattgt gacttttact	60
gttatattaa ttgccaatat aagtaaatat agattatata tgtatagtgt ttcacaaagc	120
ttagaccttt accttccagc caccacacag tgcttgatat ttcagagtca gtcattgggt	180
atacatgtgt agttccaaag cacataagct agaanaanaa atatttctag ggagcactac	240
catctgtttt cacatgaaat gccacacaca tagaactcca acatcaattt cattgcacag	300
a	301

&lt;210&gt; 280

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 280

ggtactggag ttttcctccc ctgtgaaaac gtaactactg ttgggagtga attgaggatg	60
tagaaagggtg gtggaaccaa attgtggta atggaaatag gagaatatgg ttctcactct	120
tgagaaaaaa acctaaagatt agcccaggtg gttgcctgta acttcagttt ttctgcctgg	180
gtttgatata gtttaggggtt ggggttagat taagatctaa attacatcag gacaaagaga	240
cagactatta actccacagt taattaagga ggtatgttcc atgtttattt gttaaagcag	300
t	301

&lt;210&gt; 281

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 281

aggtacaaga aggggaatgg gaaagagctg ctgctgtggc attgttcaac ttggatattc	60
gccgagcaat ccaaatcctg aatgaagggg catcttctga aaaaggagat ctgaatctca	120
atgtggtagc aatggcttta tcgggttata cggtagagaa gaactccctt tggagagaaa	180
tgtgtagcac actgcgatta cagctaaata acccgattt gtgtgtcatg tttgcatttc	240

tgacaagtga aacaggatct tacgatggag ttttgtatga aaacaaagtt gcagtacctc 300  
g 301

<210> 282  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 282  
caggtactac agaattaaaa tactgacaag caagtagttt cttggcgtgc acgaattgca 60  
tccagaacct aaaaattaag aaattcaaaa agacattttg tgggcacctg ctagcacaga 120  
agcgcagaag caaagcccag gcagaacctat gctaacctta cagctcagcc tgcacagaag 180  
cgcagaagca aagcccaggc agaacctatg taaccttaca gctcagcctg cacagaagcg 240  
cagaagcaaa gccaggcag aacatgctaa ccttacagct cagcctgcac agaagcacag 300  
a 301

<210> 283  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 283  
atctgtatac ggcagacaaa ctttatarag tgtagagagg tgagcgaaag gatgcaaaag 60  
cactttgagg gctttataat aatatgctgc ttgaaaaaaa aaatgtgtag ttgatactca 120  
gtgcatctcc agacatagta aggggttgc ctgaccaatc aggtgatcat ttttctatc 180  
acttcccagg ttttatgcaa aaattttgtt aaattctata atggtgatat gcatctttta 240  
ggaaacatat acatttttaa aaatctattt tatgtaagaa ctgacagacg aatttgcttt 300  
g 301

<210> 284  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 284  
caggtacaaa acgctattaa gtggcttaga atttgaacat ttgtggtctt tatttacttt 60  
gcttcgtgtg tgggcaaagc aacatcttcc ctaaataat attaccaaga aaagcaagaa 120  
gcagattagg tttttgacaa aacaaacagg ccaaaagggg gctgacctgg agcagagcat 180  
ggtgagaggc aaggcatgag agggcaagtt tggttgaggc agatctgtgc ctactttatt 240  
actggagtaa aagaaaacaa agttcattga tgtcgaagga tatatacagt gttagaatt 300  
a 301

<210> 285  
<211> 301  
<212> DNA  
<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(301)

<223> n = A,T,C or G.

<400> 285  
acatcaccat gatcggatcc cccacccatt atacgttgta tgtttacata aatactcttc 60  
aatgatcatt agtgttttaa aaaaaatact gaaaactcct tctgcatccc aatctctaac 120

caggaaagca aatgctatctt acagacctgc aagccctccc tcaaacnaaa ctatctctgg 180  
attaaatatt tctgacttct tttgaggtca cagcactagg caaatgctat ttacgatctg 240  
caaaagctgt ttgaagagtc aaagccccc tgtgaacacg atttctggac cctgtaacag 300  
t 301

<210> 286  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 286  
taccactgca ttccagcctg ggtgacagag tgagactccg tctccaaaaa aaactttgct 60  
tgtatattat ttttgcctta cagtggatca ttctagtagg aaaggacagt aagatttttt 120  
atcaaaatgt gtcatgccag taagagatgt tatattcttt tctcatttct tccccaccca 180  
aaaataagct accatatagc ttataagtct caaatttttg ccttttacta aaatgtgatt 240  
gtttctgttc attgtgtatg ctccatcacc tatattaggg aaattccatt ttttccttg 300  
t 301

<210> 287  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 287  
tacagatctg ggaactaaat attaaaaatg agtgtggctg gatatatgga gaatgttggg 60  
cccagaagga acgtagagat cagatattac aacagctttg ttttgagggg tagaaatatt 120  
aaatgatttg gttatgaacg cacagtttag gcagcagggc cagaatcctg accctctgcc 180  
ccgtgggtat ctctctccca gcttggctgc ctcatgttat cacagtattc cattttgttt 240  
gttgcatgtc ttgtgaagcc atcaagattt tctcgtctgt tttcctctca ttggtaatgc 300  
t 301

<210> 288  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 288  
gtacacctaa ctgcaaggac agctgaggaa tgtaatgggc agccgctttt aaagaagtag 60  
agtcaatagg aagacaaatt ccagttccag ctcatctggg gtatctgcaa agctgcaaaa 120  
gatcttttaa gacaatttca agagaatatt tccttaaagt tggcaatttg gagatcatac 180  
aaaagcatct gcttttgtga tttaatttag ctcatctggc cactggaaga atccaaacag 240  
tctgccttaa ttttggatga atgcatgatg gaaattcaat aatttagaaa gttaaaaaaa 300  
a 301

<210> 289  
<211> 301  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(301)  
<223> n = A,T,C or G

<400> 289

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ggtacactgt ttccatgta tgtttctaca cattgctacc tcagtgtcc tggaaactta      60
gcttttgatg tctccaagta gtccaccttc atttaactct ttgaaactgt atcatctttg      120
ccaagtaaga gtggtggcct atttcagctg ctttgacaaa atgactggct cctgacttaa      180
cgttctataa atgaatgtgc tgaagcaaag tgcccatggg ggcggcgaan aagagaaaga      240
tgtgttttgt tttggactct ctgtgggtcc ttccaatgct gtgggtttcc aaccagnnga      300
a                                                                 301

```

&lt;210&gt; 290

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 290

```

acactgagct cttcttgata aatatacaga atgcttggca tatacaagat tctatactac      60
tgactgatct gtccatttct ctcacagctc ttaccccaa aagcttttcc accctaagtg      120
ttctgacctc ctttttctaat cacagtaggg atagaggcag anccacctac aatgaacatg      180
gagttctatc aagaggcaga aacagcacag aatcccagtt ttaccattcg ctagcagtgc      240
tgccttgaac aaaaacattt ctccatgtct cattttcttc atgcctcaag taacagtgag      300
a                                                                 301

```

&lt;210&gt; 291

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 291

```

caggtaacaa tttcttctat cctagaaaca ttctatttta tgttggtgaa acataacaac      60
tatacagct agattttttt tctatgcttt acctgctatg gaaaatttga cacattctgc      120
tttactcttt tgtttatagg tgaatcacia aatgtatttt tatgtattct gtagttcaat      180
agccatggct gtttacttca ttttaattat ttagcataaa gacattatga aaaggcctaa      240
acatgagctt cacttcccca ctaactaatt agcatctggt atttcttaac cgtaatgcct      300
a                                                                 301

```

&lt;210&gt; 292

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 292

```

accttttagt agtaatgtct aataataaat aagaaatcaa ttttataagg tccatatagc      60
tgtattaaat aatttttaag tttaaaagat aaaataccat cattttaa atgttggtattc      120
aaaaccaaag natataaccg aaaggaaaaa cagatgagac ataaaatgat ttgcnagatg      180
ggaaatatag tasttyatga atgttnatta aattccagtt ataatagtgg ctacacactc      240
tcactacaca cacagacccc acagtcctat atgccacaaa cacatttcca taacttgaaa      300
a                                                                 301

```

<210> 293  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 293  
 ggtaccaagt gctgggtgcca gcctgttacc tgttctcact gaaaagtctg gctaattgctc 60  
 ttgtgtagtc acttctgatt ctgacaatca atcaatcaat ggcctagagc actgactgtt 120  
 aacacaaaacg tctactagcaa agtagcaaca gctttaagtc taaatacaaa gctgttctgt 180  
 gtgagaattt tttaaaaggc tacttgtata ataacccttg tcatttttaa tgtacctcgg 240  
 ccgcgaccac gctaagccga attctgcaga tatccatcac actggcgggc gctcgagcat 300  
 g 301

<210> 294  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<220> -----  
 <221> misc\_feature  
 <222> (1)...(301)  
 <223> n = A,T,C or G

<400> 294  
 tgaccataaa caatatacac tagctatctt tttactgtc catcattagc accaatgaag 60  
 attcaataaa attaccttta ttcacacatc tcaaaacaat tctgcaaatt cttagtgaag 120  
 tttactata gtcacaganc ttaaatattc acattgtttt ctatgtctac tgaaaataag 180  
 ttcactactt ttctgggata ttctttacaa aatcttatta aaattcctgg tattatcacc 240  
 cccaattata cagtagcaca accaccttat gtagttttta catgatagct ctgtagaggt 300  
 t 301

<210> 295  
 <211> 305  
 <212> DNA  
 <213> Homo sapien

<400> 295  
 gtactctttc tctcccttcc tctgaattta attctttcaa cttgcaattt gcaaggatta 60  
 cacatttcac tgtgatgtat attgtgttgc aaaaaaaaaa gtgtctttgt ttaaaattac 120  
 ttggtttttg aatccatctt gctttttccc cattggaact agtcattaac ccatctctga 180  
 actggtagaa aaacrtctga agagctagtc tatcagcatc tgacaggtga attggatggt 240  
 tctcagaacc atttcaccca gacagcctgt ttctatcctg tttataaat tagtttgggt 300  
 tctct 305

<210> 296  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 296  
 aggtactatg ggaagctgct aaaataatat ttgatagtaa aagtatgtaa tgtgctatct 60  
 cacctagtag taaactaaaa ataaactgaa actttatgga atctgaagtt attttccttg 120  
 attaaataga attaataaac caatatgagg aaacatgaaa ccatgcaatc tactatcaac 180  
 tttgaaaaag tgattgaacg aaccacttag ctttcagatg atgaacactg ataagtcatt 240



tgtcattact ataaatttta aaatctgtta ataagatggc ctatagggag gaaaaagggg 300  
c 301

<210> 297

<211> 300

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(300)

<223> n = A,T,C or G

<400> 297

actgagtttt aactggacgc caagcaggca aggctggaag gttttgctct ctttggtgcta 60  
aaggttttga aaaccttgaa ggagaatcat tttgacaaga agtacttaag agtctagaga 120  
acaaagangt gaaccagctg aaagctctcg ggggaanctt acatgtgttg ttaggcctgt 180  
tccatcattg ggagtgcact ggccatccct caaaatttgt ctgggctggc ctgagtggtc 240  
accgcacctc ggccgcgacc acgctaagcc gaattctgca gatatccatc acactggcgg 300

<210> 298

<211> 301

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(301)

<223> n = A,T,C or G

<400> 298

tatggggttt gtcacccaaa agctgatgct gagaaaggcc tccctggggc ccctcccgcg 60  
ggcatctgag agacctggtg ttccagtgtt tctggaaatg ggtcccagtg ccgcccggctg 120  
tgaagctctc agatcaatca cgggaagggc ctggcggtgg tggccacctg gaaccaccct 180  
gtcctgtctg tttacatttc actaycaggt tttctctggg cattacnatt tgttccccta 240  
caacagtgc ctgtgcattc tgctgtggcc tgctgtgtct gcaggtggct ctcagcgagg 300  
t 301

<210> 299

<211> 301

<212> DNA

<213> Homo sapien

<400> 299

gttttgagac ggagtttcac tcttggtgcc cagactggac tgcaatggca gggctctctgc 60  
tactgcacc ctctgcctcc caggttcgag caattctcct gcctcagcct cccaggtagc 120  
tgggattgca ggctcacgcc accataccca gctaattttt ttgtattttt agtagagacg 180  
gagtttcgcc atgttggccg gctggtctca aactcctgac ctcaagcgac ctgcctgcct 240  
cggcctccca aagtgtctgga attataggca tgagtcaaca cgcccagcct aaagatatatt 300  
t 301

<210> 300

<211> 301

<212> DNA

<213> Homo sapien

&lt;400&gt; 300

attcagtttt	atttgctgcc	ccagtatctg	taaccaggag	tgccacaaaa	tcttgccaga	60
tatgtccac	acccactggg	aaaggctccc	acctggctac	ttcctctatc	agctgggtca	120
gctgcattcc	acaaggttct	cagcctaata	agtttacta	cctgccagtc	tcaaaactta	180
gtaaagcaag	accatgacat	tccccacgg	aaatcagagt	ttgccccacc	gtcttgttac	240
tataaagcct	gcctctaaca	gtccttgctt	cttcacacca	atccccgagcg	catcccccat	300
g						301

&lt;210&gt; 301

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 301

ttaaattttt	gagaggataa	aaaggacaaa	taatctagaa	atgtgtcttc	ttcagtctgc	60
agaggacccc	aggtctccaa	gcaaccacat	ggtcaagggc	atgaataatt	aaaagttggt	120
gggaactcac	aaagaccctc	agagctgaga	caccacacac	agtgggagct	cacaaagacc	180
ctcagagctg	agacaccac	aacagtggga	gtcacaag	accctcagag	ctgagacacc	240
cacaacagca	cctcgttcag	ctgccacatg	tgtgaataag	gatgcaatgt	ccagaagtgt	300
t						301

&lt;210&gt; 302

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 302

aggtacacat	ttagcttggtg	gtaaatgact	cacaaaactg	attttaaaat	caagttaatg	60
tgaattttga	aaattactac	ttaatcctaa	ttcacataa	caatggcatt	aaggtttgac	120
ttgagttggt	tcttagtatt	atttatggta	aataggctct	taccacttgc	aaataactgg	180
ccacatcatt	aatgactgac	ttcccagtaa	ggctctctaa	ggggtaagta	ggaggatcca	240
caggatttga	gatgctaagg	ccccagagat	cgtttgatcc	aaccctctta	ttttcagagg	300
g						301

&lt;210&gt; 303

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 303

aggtaccaac	tgtggaaata	ggtagaggat	cattttttct	ttccatatca	actaagttgt	60
atattgtttt	ttgacagttt	aacacatctt	cttctgtcag	agattctttc	acaatagcac	120
tggctaatag	aactaccgct	tgcattgtaa	aaatgggtgt	ttgtgaaatg	atcataggcc	180
agtaacgggt	atgtttttct	aactgatctt	ttgctcgttc	caaagggacc	tcaagacttc	240
catcgatttt	atatctgggg	tctagaaaag	gagttaatct	gttttccctc	ataaattcac	300
c						301

&lt;210&gt; 304

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 304

acatggatgt	tattttgcag	actgtcaacc	tgaatttgta	tttgcttgac	attgcctaata	60
------------	------------	------------	------------	------------	-------------	----

tattagtttc agtttcagct taccacttt ttgtctgcaa catgcaraas agacagtgcc	120
cttttttagtg tatcatatca ggaatcatct cacattgggt tgtgccatta ctgggtgcagt	180
gactttcagc cacttgggta aggtggagtt ggccatatgt ctccactgca aaattactga	240
ttttcctttt gtaattaata agtgtgtgtg tgaagattct ttgagatgag gtatatatct	300
c	301

&lt;210&gt; 305

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 305

gangtacagc gtggtcaagg taacaagaag aaaaaaatgt gagtggcatc ctgggatgag	60
caggggggaca gacctggaca gacacgttgt catttgctgc tgtgggtagg aaaatgggag	120
taaaggagga gaaacagata caaaatctcc aactcagtat taaggatttc tcatgcctag	180
aatattggta gaaacaagaa tacattcata tggcaaataa ctaaccatgg tggacaacaaa	240
ttctgggatt taagttggat accaangaaa ttgtattaaa agagctgttc atggaataag	300
a	301

&lt;210&gt; 306

&lt;211&gt; 8

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 306

Val Leu Gly Trp Val Ala Glu Leu

1

5

&lt;210&gt; 307

&lt;211&gt; 637

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 307

acagggratg aagggaagg gagaggatga ggaagcccc ctggggattt ggtttggtcc	60
ttgtgatcag gtggtctatg gggcttatcc ctacaaagaa gaatccagaa atagggggcac	120
attgaggaat gatacttgag cccaaagagc attcaatcat tgttttattt gccttmtttt	180
cacaccattg gtgagggagg gattaccacc ctggggttat gaagatggtt gaacacccca	240
cacatagcac cggagatatg agatcaacag tttcttagcc atagagattc acagcccaga	300
gcaggaggac gcttgcacac catgcaggat gacatggggg atgcgctcgg gattggtgtg	360
aagaagcaag gactgttaga ggcaggcttt atagtaacaa gacggtgggg caaactctga	420
tttccgtggg ggaatgtcat ggtcttgctt tactaagttt tgagactggc aggtagtga	480
actcattagg ctgagaacct tgtggaatgc acttgaccca sctgatagag gaagtagcca	540
gggtgggagcc tttccagtg ggtgtgggac atatctggca agattttgtg gcactcctgg	600
ttacagatac tggggcagca aataaaactg aatcttg	637

&lt;210&gt; 308

&lt;211&gt; 647

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(647)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 308

acgattttca	ttatcatgta	aatcgggtca	ctcaaggggc	caaccacagc	tgggagccac	60
tgctcagggg	aaggttcata	tgggactttc	tactgcccac	ggttctatac	aggatataaa	120
ggngcctcac	agtatagatc	tggtagcaaa	gaagaagaaa	caaacactga	tctctttctg	180
ccacccctct	gacccttttg	aactcctctg	accctttaga	acaagcctac	ctaatatctg	240
ctagagaaaa	gaccaacaac	ggcctcaaag	gatctcttac	catgaaggtc	tcagctaatt	300
cttggctaag	atgtgggttc	cacattaggt	tctgaatatg	gggggaaggg	tcaatttgct	360
cattttgtgt	gtggataaa	tcaggatgcc	cagggggccag	agcagggggc	tgcttgcttt	420
gggaacaatg	gctgagcata	taacataggt	ttatggggaa	caaaacaaca	tcaaagtcac	480
tgtatcaatt	gccatgaaga	cttgagggac	ctgaatctac	cgattcatct	taaggcagca	540
ggaccagttt	gagtggcaac	aatgcagcag	cagaatcaat	ggaaacaaca	gaatgattgc	600
aatgtccttt	ttttctcct	gcttctgact	tgataaaaag	ggaccgt		647

&lt;210&gt; 309

&lt;211&gt; 460

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 309

actttatagt	ttaggctgga	cattggaaaa	aaaaaaaaagc	cagaacaaca	tgtgatagat	60
aatatgattg	gctgcacact	tccagactga	tgaatgatga	acgtgatgga	ctattgtatg	120
gagcacatct	tcagcaagag	ggggaaatac	tcatcatttt	tggccagcag	ttgtttgatc	180
accaaacatc	atgccagaat	actcagcaaa	ccttcttagc	tcttgagaag	tcaaagtccg	240
ggggaattta	ttcctggcaa	ttttaattgg	actccttatg	tgagagcagc	ggctaccag	300
ctggggtggt	ggagcgaacc	cgtcactagt	ggacatgcag	tggcagagct	cctggtaacc	360
acctagagga	atacacaggc	acatgtgtga	tgccaagcgt	gacacctgta	gcactcaaat	420
ttgtcttggt	tttgtctttc	ggtgtgtaag	attcttaagt			460

&lt;210&gt; 310

&lt;211&gt; 539

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 310

acgggactta	tcaaataaag	ataggaaaag	aagaaaactc	aaatattata	ggcagaaatg	60
ctaaaggttt	taaaatatgt	caggattgga	agaaggcatg	gataaagaac	aaagttcagt	120
taggaaagag	aaacacagaa	ggaagagaca	caataaaaag	cattatgtat	tctgtgagaa	180
gtcagacagt	aagatttggt	ggaaatgggt	tggtttggtg	tatgggtatg	attttagcaa	240
taatctttat	ggcagagaaa	gctaaaatcc	tttagcttgc	gtgaatgatc	acttgctgaa	300
ttcctcaagg	taggcatgat	gaaggagggt	ttagaggaga	cacagacaca	atgaactgac	360
ctagatagaa	agccttagta	tactcagcta	ggaatagtga	ttctgagggc	acactgtgac	420
atgattatgt	cattacatgt	atggtagtga	tggggatgat	aggaaggaag	aacttatggc	480
atattttcac	ccccacaaa	gtcagttaaa	tattgggaca	ctaaccatcc	aggtcaaga	539

&lt;210&gt; 311

&lt;211&gt; 526

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(526)  
 <223> n = A,T,C or G

<400> 311  
 caaatttgag ccaatgacat agaattttac aaatcaagaa gcttattctg gggccatttc 60  
 ttttgacgtt ttctctaaac tactaaagag gcattaatga tccataaatt atattatcta 120  
 catttacagc atttaaaatg tggtcagcat gaaatattag ctacagggga agctaaataa 180  
 attaaacatg gaataaagat ttgtccttaa atataatcta caagaagact ttgatatttg 240  
 tttttcacaa gtgaagcatt cttataaagt gtcataacct ttttggggaa actatgggaa 300  
 aaaatgggga aactctgaag ggttttaagt atcttacctg aagctacaga ctccataacc 360  
 tctctttaca gggagctcct gcagcccta cagaaatgag tggctgagat tcttgattgc 420  
 acagcaagag cttctcatct aaaccctttc cctttttagt atctgtgtat caagtataaa 480  
 agttctataa actgtagtnt acttatttta atccccaaag cacagt 526

<210> 312  
 <211> 500  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(500)  
 <223> n = A,T,C or G

<400> 312  
 cctctctctc cccacccct gactctagag aactgggtt tctcccagta ctccagcaat 60  
 tcatttctga aagcagttga gccactttat tccaaagtag actgcagatg ttcaaactct 120  
 ccatttctct ttccttcca cctgccagtt ttgctgactc tcaacttgtc atgagtgtaa 180  
 gcattaagga cattatgctt cttcgattct gaagacaggc cctgctcatg gatgactctg 240  
 gcttcttagg aaaatatttt tcttccaaaa tcagtaggaa atctaaactt atccccctt 300  
 tgcagatgct tagcagcttc agacatttgg ttaagaaccc atgggaaaaa aaaaaatcct 360  
 tgctaatttg gtttcttttg taaaccanga ttcttatttg nctggtatag aatatcagct 420  
 ctgaacgtgt ggtaaagatt tttgtgtttg aatataggag aaatcagttt gctgaaaagt 480  
 tagtcttaat tatctattgg 500

<210> 313  
 <211> 718  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(718)  
 <223> n = A,T,C or G

<400> 313  
 ggagatttgt gtggtttgca gccgagggag accaggaaga tctgcatggt gggaaggacc 60  
 tgatgataca gaggtgagaa ataagaaagg ctgctgactt taccatctga ggccacacat 120  
 ctgctgaaat ggagataatt aacatcacta gaaacagcaa gatgacaata taatgtctaa 180  
 gtagtgacat gtttttgac atttccagcc cttttaaata tccacacaca caggaagcac 240  
 aaaaggaagc acagagatcc ctgggagaaa tgcccgcccg ccattctggg tcatcgatga 300  
 gctcgcct gtgcctgntc ccgcttgatg ggggaaggaca ttagaaaatg aattgatgtg 360  
 ttccttaaaag gatggcagga aaacagatcc tgttgtggat atttatttga acgggattac 420

agatttgaaa tgaagtcaca aagtgagcat taccaatgag aggaaaacag acgagaaaaat	480
cttgatgggt cacaagacat gcaacaaaca aaatggaata ctgtgatgac acgagcagcc	540
aactggggag gagataccac ggggcagagg tcaggattct ggccctgctg cctaactgtg	600
cgttatacca atcattttcta tttctaccct caaacaagct gtngaataac tgacttacgg	660
ttcttntggc ccacattttc atnatccacc ccntcntttt aannttantic caaantgt	718

&lt;210&gt; 314

&lt;211&gt; 358

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 314

gtttattttac attacagaaa aaacatcaag acaatgtata ctattttcaaa tatatccata	60
cataatcaaa tatagctgta gtacatgttt tcattgggtg agattaccac aaatgcaagg	120
caacatgtgt agatctcttg tcttattctt ttgtctataa tactgtattg tgtagtccaa	180
gctctcggtg gtccagccac tgtgaaacat gctcccttta gattaacctc gtggacgctc	240
ttgttgatt gctgaactgt agtgccctgt attttgcttc tgtctgtgaa ttctgttget	300
tctggggcat ttccttgatga tgcagaggac caccacacag atgacagcaa tctgaatt	358

&lt;210&gt; 315

&lt;211&gt; 341

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 315

taccacctcc ccgctggcac tgatgagccg catcaccatg gtcaccagca ccatgaaggc	60
ataggtgatg atgaggacat ggaatgggcc cccaaggatg gtctgtccaa agaagcgagt	120
gacccccatt ctgaagatgt ctggaacctc taccagcagg atgatgatag cccaatgac	180
agtcaccagc tccccgacca gccggatata gtccttaggg gtcattgtag cttcctgaag	240
tagcttctgc tgtaagaggg tggtgtcccg ggggctcgtg cggttattgg tcttgggctt	300
gagggggcg tagatgcagc acatgggtgaa gcagatgatg t	341

&lt;210&gt; 316

&lt;211&gt; 151

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 316

agactgggca agactcttac gcccacact gcaatttggc cttgttgccg tatccattta	60
tgtgggcctt tctcgagttt ctgattataa acaccactgg agcgatgtgt tgactggact	120
cattcagggg gctctggttg caatattagt t	151

&lt;210&gt; 317

&lt;211&gt; 151

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 317

agaactagtg gatcctaagt aaataacctga aacatatatt ggcatttatc aatgggtcaa	60
atcttcattt atctctggcc ttaaccctgg ctctgaggc tgcggccagc agatcccagg	120
ccagggctct gttcttgcca cacctgcttg a	151

&lt;210&gt; 318

&lt;211&gt; 151

&lt;212&gt; DNA

<213> Homo sapien

<400> 318

```
actggtggga ggcgctgttt agttggctgt ttccagaggg gtctttcgga gggacctcct    60
gctgcaggct ggagtgtctt tttcctggc gggagaccgc acattccact gctgaggctg    120
tgggggcggt ttatcaggca gtgataaaca t                                151
```

<210> 319

<211> 151

<212> DNA

<213> Homo sapien

<400> 319

```
aactagtggga tccagagcta taggtacagt gtgatctcag ctttgcaaac acattttcta    60
catagatagt actaggtatt aatagatatg taaagaaaga aatcacacca ttaataatgg    120
taagattggg tttatgtgat tttagtggg a                                151
```

<210> 320

<211> 150

<212> DNA

<213> Homo sapien

<400> 320

```
aactagtggga tccactagtc cagtgtgggtg gaattccatt gtgttgggggt tctagatcgc    60
gagcggctgc cttttttttt ttttttttg ggggggaatt tttttttttt aatagttatt    120
gagtgttcta cagcttacag taaataccat                                151
```

<210> 321

<211> 151

<212> DNA

<213> Homo sapien

<400> 321

```
agcaactttg tttttcatcc aggttatattt aggccttagga ttccctctca cactgcagtt    60
tagggtgga ttgtaaccag ctatggcata ggtgttaacc aaaggctgag taaacatggg    120
tgcctctgag aatcaaagt cttcatacac t                                151
```

<210> 322

<211> 151

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(151)

<223> n = A,T,C or G

<400> 322

```
atccagcacc ttctcctgtt tttgccttc cttttcttc ttcttasatt ctgcttgagg    60
tttgggcttg gtcagtttgc cacagggctt ggagatgggt acagtcttct ggcattcggc    120
attgtgcagg gctcgttca nacttccagt t                                151
```

<210> 323

<211> 151

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(151)

<223> n = A,T,C or G

<400> 323

tgaggacttg tktttttttt ctttattttt aatcctctta ckttgtaa	atattgccta	60
nagactcant tactaccag tttgtggtt	ttgtggagaa atgtaactgg acagttagct	120
gttcaatyaa aaagacatt ancccatgtg	g	151

<210> 324

<211> 461

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(461)

<223> n = A,T,C or G

<400> 324

acctgtgtgg aatttcagct ttcctcatgc aaaaggattt	tgtatccccg gcctacttga	60
agaagtggc agctaaagga atccagggtg ttggttgac	tgtaataacc tttgatgaaa	120
agagttacta cgaatcccat cttgggtcca gctatatcac	tgacagcatg gtagaagact	180
gcgaacctca cttctagact ttcacgggtg gacgaaacgg	gttcagaaac tgcagggggc	240
ctcatacagg gatatcaaaa taccctttgt gctaccagg	cctgggggaa tcagggtgact	300
cacacaaatg caatagttg tcaactgcatt tttacctgaa	ccaaagctaa acccggtgtt	360
gccaccatgc accatggcat gccagagttc aacactgttg	ctcttgaaaa ttgggtctga	420
aaaaacgcac aagagcccct gccctgcct agctgangca	c	461

<210> 325

<211> 400

<212> DNA

<213> Homo sapien

<400> 325

acactgtttc catgttatgt ttctacacat tgctacctca	gtgctcctgg aaacttagct	60
tttgatgtct ccaagtagtc caccttcatt taactcttg	aaactgtatc atctttgcca	120
agtaagagtg gtggcctatt tcagctgctt tgacaaaatg	actggctcct gacttaacgt	180
tctataaatg aatgtgctga agcaaagtgc ccatggtggc	ggcgaagaag agaaagatgt	240
gttttgttt ggactctctg tggctccttc caatgctgtg	ggtttccaac cagggaagg	300
gtccctttt cattgccaag tgccataacc atgagcacta	cgctaccatg gttctgcctc	360
ctggccaagc aggtgtgtt gcaagaatga aatgaatgat		400

<210> 326

<211> 1215

<212> DNA

<213> Homo sapien

<400> 326

ggaggactgc agcccgact cgcagccctg gcaggcgga	ctggctcatgg aaaacgaatt	60
gttctgctcg ggcgtcctgg tgcacccgca gtgggtgctg	tcagccgcac actgtttcca	120
gaactcctac accatcgggc tgggcctgca cagtcttgag	gccgaccaag agccaggag	180



ccagatggtg	gaggccagcc	tctccgtacg	gcacccagag	tacaacagac	ccttgctcgc	240
taacgacctc	atgctcatca	agttggacga	atccgtgtcc	gagtcctgaca	ccatccggag	300
catcagcatt	gcttcgcagt	gccctaccgc	ggggaactct	tgcctcgttt	ctggctgggg	360
tctgctggcg	aacggcagaa	tgcctaccgt	gctgcagtgc	gtgaacgtgt	cggtggtgtc	420
tgaggaggtc	tgcagtaagc	tctatgaccc	gctgtaccac	cccagcatgt	tctgcgccgg	480
cggagggcaa	gaccagaagg	actcctgcaa	cggtgactct	ggggggcccc	tgatctgcaa	540
cgggtacttg	cagggccttg	tgtctttcgg	aaaagccccg	tgtggccaag	ttggcggtgcc	600
aggtgtctac	accaacctct	gcaaattcac	tgagtggata	gagaaaaccg	tccaggccag	660
ttaactctgg	ggactgggaa	cccatgaaat	tgaccccaa	atacatcctg	cggaaggaat	720
tcaggaatat	ctgttcccag	ccccctctcc	ctcaggccca	ggagtccagg	ccccagcccc	780
ctcctccctc	aaaccaaggg	tacagatccc	cagccccctc	tccctcagac	ccaggagtcc	840
agacccccca	gccccctctc	cctcagaccc	aggagtccag	ccccctctcc	ctcagaccca	900
ggagtccaga	ccccccagcc	cctcctccct	cagacccagg	ggtccaggcc	cccaaccctt	960
cctccctcag	actcagaggt	ccaagccccc	aaccctctct	tccccagacc	cagaggtcca	1020
ggtcccagcc	cctcctccct	cagacccagc	ggtccaatgc	cacctagact	ctccctgtac	1080
acagtgcccc	cttgtggcac	gttgacccaa	ccttaccagt	tggtttttca	ttttttgtcc	1140
ctttccccta	gatccagaaa	taaagtctaa	gagaagcgca	aaaaaaaaaa	aaaaaaaaaa	1200
aaaaaaaaaa	aaaaaa					1215

&lt;210&gt; 327

&lt;211&gt; 220

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 327

Glu	Asp	Cys	Ser	Pro	His	Ser	Gln	Pro	Trp	Gln	Ala	Ala	Leu	Val	Met
1				5				10						15	
Glu	Asn	Glu	Leu	Phe	Cys	Ser	Gly	Val	Leu	Val	His	Pro	Gln	Trp	Val
			20					25					30		
Leu	Ser	Ala	Ala	His	Cys	Phe	Gln	Asn	Ser	Tyr	Thr	Ile	Gly	Leu	Gly
			35				40					45			
Leu	His	Ser	Leu	Glu	Ala	Asp	Gln	Glu	Pro	Gly	Ser	Gln	Met	Val	Glu
	50				55					60					
Ala	Ser	Leu	Ser	Val	Arg	His	Pro	Glu	Tyr	Asn	Arg	Pro	Leu	Leu	Ala
65					70					75					80
Asn	Asp	Leu	Met	Leu	Ile	Lys	Leu	Asp	Glu	Ser	Val	Ser	Glu	Ser	Asp
			85					90					95		
Thr	Ile	Arg	Ser	Ile	Ser	Ile	Ala	Ser	Gln	Cys	Pro	Thr	Ala	Gly	Asn
			100					105					110		
Ser	Cys	Leu	Val	Ser	Gly	Trp	Gly	Leu	Leu	Ala	Asn	Gly	Arg	Met	Pro
			115				120					125			
Thr	Val	Leu	Gln	Cys	Val	Asn	Val	Ser	Val	Val	Ser	Glu	Glu	Val	Cys
			130			135					140				
Ser	Lys	Leu	Tyr	Asp	Pro	Leu	Tyr	His	Pro	Ser	Met	Phe	Cys	Ala	Gly
145					150					155					160
Gly	Gly	Gln	Asp	Gln	Lys	Asp	Ser	Cys	Asn	Gly	Asp	Ser	Gly	Gly	Pro
			165					170					175		
Leu	Ile	Cys	Asn	Gly	Tyr	Leu	Gln	Gly	Leu	Val	Ser	Phe	Gly	Lys	Ala
			180				185						190		
Pro	Cys	Gly	Gln	Val	Gly	Val	Pro	Gly	Val	Tyr	Thr	Asn	Leu	Cys	Lys
			195				200					205			
Phe	Thr	Glu	Trp	Ile	Glu	Lys	Thr	Val	Gln	Ala	Ser				
	210					215					220				

&lt;210&gt; 328

&lt;211&gt; 234

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 328

cgctcgtctc	tggtagctgc	agccaaatca	taaacggcga	ggactgcagc	ccgcactcgc	60
agccctggca	ggcggcactg	gtcatggaaa	acgaattgtt	ctgctcgggc	gtcctgggtgc	120
atccgcagtg	ggtgctgtca	gccacacact	gtttccagaa	ctcctacacc	atcgggctgg	180
gcctgcacag	tcttgaggcc	gaccaagagc	cagggagcca	gatgggtggag	gcca	234

&lt;210&gt; 329

&lt;211&gt; 77

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 329

Leu	Val	Ser	Gly	Ser	Cys	Ser	Gln	Ile	Ile	Asn	Gly	Glu	Asp	Cys	Ser	1	5	10	15
Pro	His	Ser	Gln	Pro	Trp	Gln	Ala	Ala	Leu	Val	Met	Glu	Asn	Glu	Leu	20	25	30	35
Phe	Cys	Ser	Gly	Val	Leu	Val	His	Pro	Gln	Trp	Val	Leu	Ser	Ala	Thr	35	40	45	50
His	Cys	Phe	Gln	Asn	Ser	Tyr	Thr	Ile	Gly	Leu	Gly	Leu	His	Ser	Leu	50	55	60	65
Glu	Ala	Asp	Gln	Glu	Pro	Gly	Ser	Gln	Met	Val	Glu	Ala				70	75		

&lt;210&gt; 330

&lt;211&gt; 70

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 330

cccaacacaa	tggcccgatc	ccatccctga	ctccgccctc	aggatcgctc	gtctctggta	60
gctgcagcca						70

&lt;210&gt; 331

&lt;211&gt; 22

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 331

Gln	His	Asn	Gly	Pro	Ile	Pro	Ser	Leu	Thr	Pro	Pro	Ser	Gly	Ser	Leu	1	5	10	15
Val	Ser	Gly	Ser	Cys	Ser											20			

&lt;210&gt; 332

&lt;211&gt; 2507

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 332

tggtgccgct	gcagccggca	gagatgggtg	agctcatgtt	cccgtgttg	ctctccttc	60
tgcccttct	tctgtatatg	gctgcgcccc	aaatcaggaa	aatgctgtcc	agtgggggtg	120

gtacatcaac	tgttcagctt	cctgggaaaag	tagttgtggt	cacaggagct	aatacaggta	180
tcgggaagga	gacagccaaa	gagctggctc	agagaggagc	tcgagtatat	ttagcttgcc	240
gggatgtgga	aaagggggaa	ttggtggcca	aagagatcca	gaccacgaca	gggaaccagc	300
aggtgttggt	gcggaaactg	gacctgtctg	atactaagtc	tattcgagct	tttgctaagg	360
gcttcttagc	tgaggaaaag	cacctccacg	ttttgatcaa	caatgcagga	gtgatgatgt	420
gtccgtactc	gaagacagca	gatggctttg	agatgcacat	aggagtcaac	cacttgggtc	480
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&lt;210&gt; 333

&lt;211&gt; 3030

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 333

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&lt;210&gt; 334

&lt;211&gt; 2417

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 334

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&lt;210&gt; 335

&lt;211&gt; 2984

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 335

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&lt;210&gt; 336

&lt;211&gt; 147

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 336

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Leu	Asp	Ser	Glu	Asn	Thr	Ser	Gly	Ala	Leu	Pro	Arg	Leu	Pro	Gln	Thr
			20					25					30		
Pro	Lys	Gln	Pro	Gln	Lys	Arg	Ser	Arg	Ala	Ala	Phe	Ser	His	Thr	Gln
		35					40					45			
Val	Ile	Glu	Leu	Glu	Arg	Lys	Phe	Ser	His	Gln	Lys	Tyr	Leu	Ser	Ala
	50					55					60				
Pro	Glu	Arg	Ala	His	Leu	Ala	Lys	Asn	Leu	Lys	Leu	Thr	Glu	Thr	Gln
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Val Lys Ile Trp Phe Gln Asn Arg Arg Tyr Lys Thr Lys Arg Lys Gln  
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 Leu Ser Ser Glu Leu Gly Asp Leu Glu Lys His Ser Ser Leu Pro Ala  
                     100                    105                    110  
 Leu Lys Glu Glu Ala Phe Ser Arg Ala Ser Leu Val Ser Val Tyr Asn  
                     115                    120                    125  
 Ser Tyr Pro Tyr Tyr Pro Tyr Leu Tyr Cys Val Gly Ser Trp Ser Pro  
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 Ala Phe Trp  
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 <212> PRT  
 <213> Homo sapien

<400> 339  
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                     20                    25                    30  
 Cys Thr Ser Thr Val Gln Leu Pro Gly Lys Val Val Val Val Thr Gly  
                     35                    40                    45  
 Ala Asn Thr Gly Ile Gly Lys Glu Thr Ala Lys Glu Leu Ala Gln Arg  
                     50                    55                    60  
 Gly Ala Arg Val Tyr Leu Ala Cys Arg Asp Val Glu Lys Gly Glu Leu  
   65                    70                    75                    80  
 Val Ala Lys Glu Ile Gln Thr Thr Thr Gly Asn Gln Gln Val Leu Val  
                     85                    90                    95  
 Arg Lys Leu Asp Leu Ser Asp Thr Lys Ser Ile Arg Ala Phe Ala Lys  
                     100                    105                    110  
 Gly Phe Leu Ala Glu Glu Lys His Leu His Val Leu Ile Asn Asn Ala  
                     115                    120                    125  
 Gly Val Met Met Cys Pro Tyr Ser Lys Thr Ala Asp Gly Phe Glu Met  
                     130                    135                    140  
 His Ile Gly Val Asn His Leu Gly His Phe Leu Leu Thr His Leu Leu

145                      150                      155                      160  
 Leu Glu Lys Leu Lys Glu Ser Ala Pro Ser Arg Ile Val Asn Val Ser  
                                  165                      170                      175  
 Ser Leu Ala His His Leu Gly Arg Ile His Phe His Asn Leu Gln Gly  
                                  180                      185                      190  
 Glu Lys Phe Tyr Asn Ala Gly Leu Ala Tyr Cys His Ser Lys Leu Ala  
                                  195                      200                      205  
 Asn Ile Leu Phe Thr Gln Glu Leu Ala Arg Arg Leu Lys Gly Ser Gly  
                                  210                      215                      220  
 Val Thr Thr Tyr Ser Val His Pro Gly Thr Val Gln Ser Glu Leu Val  
 225                                   230                                   235                                   240  
 Arg His Ser Ser Phe Met Arg Trp Met Trp Trp Leu Phe Ser Phe Phe  
                                  245                                   250                                   255  
 Ile Lys Thr Pro Gln Gln Gly Ala Gln Thr Ser Leu His Cys Ala Leu  
                                  260                                   265                                   270  
 Thr Glu Gly Leu Glu Ile Leu Ser Gly Asn His Phe Ser Asp Cys His  
                                  275                                   280                                   285  
 Val Ala Trp Val Ser Ala Gln Ala Arg Asn Glu Thr Ile Ala Arg Arg  
                                  290                                   295                                   300  
 Leu Trp Asp Val Ser Cys Asp Leu Leu Gly Leu Pro Ile Asp  
 305                                   310                                   315

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 <211> 483  
 <212> DNA  
 <213> Homo sapien

<400> 340

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 ggttggtggg gcggtttatc aggcagtgat aaacataaga tgatcatttc ttgactccgg 240  
 ccttcaattt tctctttggc tgacgacgga gtccgtggtg tcccgatgta actgaccctc 300  
 gctccaaacg tgacatcact gatgctcttc tcgggggtgc tgatggcccg cttgggtcag 360  
 tgctcaatct cgccattcga ctcttgcttc aaactgtatg aagacacctg actgcacgtt 420  
 tttctggtgc ttccagaatt taaagtgaag ggcagcactc ctaagctccg actccgatgc 480  
 ctg 483

<210> 341  
 <211> 344  
 <212> DNA  
 <213> Homo sapien

<400> 341

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 attaatataa taattttctga tgatggtttt atctgcagta atatgtatat catctattag 240  
 aatttactta atgaaaaact gaagagaaca aaatttgtaa ccactagcac ttaagtactc 300  
 ctgattctta acattgtctt taatgaccac aagacaacca acag 344

<210> 342  
 <211> 592  
 <212> DNA  
 <213> Homo sapien



## &lt;400&gt; 342

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cctggcaggt	aaaccaatgc	caagagagt	atggaaacca	ttggcaagac	tttgttgatg	180
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cccgtgtcct	tatgcaaata	atcgtcttct	tctaaatttc	tcctaggctt	cattttccaa	480
agttcttctt	ggtttgatg	gtcttttctg	ctttccatta	attctataaa	atagtatggc	540
ttcagccacc	cactcttcgc	cttagcttga	ccgtgagtct	cggctgccgc	tg	592

&lt;210&gt; 343

&lt;211&gt; 382

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

## &lt;400&gt; 343

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aaaccaccaa	gctgaaaaaa	aa				382

&lt;210&gt; 344

&lt;211&gt; 536

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

## &lt;400&gt; 344

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tcgaccctat	atcccccgcc	cgcgtccctt	tctccataaa	attcttctta	gtagctatta	360
ccttcttatt	atttgatcta	gaaattgccc	tccttttacc	cctaccatga	gccctacaaa	420
caactaacct	gccactaata	gttatgtcat	ccctcttatt	aatcatcatc	ctagccctaa	480
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&lt;210&gt; 345

&lt;211&gt; 251

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

## &lt;400&gt; 345

accttttgag	gtctctctca	ccacctccac	agccaccgtc	accgtgggat	gtgctggatg	60
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gcgtgggcca	ggaaatcaca	tcctacactg	cccaggagcc	agacacattt	atggaacaga	180
aaataacata	tcggatttgg	agagacactg	ccaactggct	ggagattaat	ccggacactg	240
gtgccatttc	c					251

<210> 346  
 <211> 282  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(282)  
 <223> n = A,T,C or G

<400> 346  
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 ctaagtcttg ttacaaaaa aaggaaaaag aaaagatctt ctcagttaca aattctggga 120  
 agggagacta tacctggctc ttgcctaag tgagaggtct tccctcccgc accaaaaaat 180  
 agaaaggctt tctatttcac tggcccaggt agggggaagg agagtaactt tgagtctgtg 240  
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<210> 347  
 <211> 201  
 <212> DNA  
 <213> Homo sapien

<220>  
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 <222> (1)...(201)  
 <223> n = A,T,C or G

<400> 347  
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 tctgagactg actggaccca ccagaccca gggcaaagat acatgttacc atatcatctt 180  
 tataaagaat ttttttttgt c 201

<210> 348  
 <211> 251  
 <212> DNA  
 <213> Homo sapien

<400> 348  
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 aggagacact cccagcatgg aggaggggtt atcttttcat cctaggtcag gtctacaatg 180  
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 gccctgcctc c 251

<210> 349  
 <211> 251  
 <212> DNA  
 <213> Homo sapien

<400> 349  
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 cagaagggtc tgaactctac gtgtaccag agaacataat gcaattcatg cattccactt 180  
 agcaattttg taaaatacca gaaacagacc ccaagagtct ttcaagatga ggaaaattca 240

actcctgggtt t

251

&lt;210&gt; 350

&lt;211&gt; 908

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 350

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cggctggaat	tgctctgggt	atgatgacag	agaaaatgat	ctcttcctct	gtgacaccaa	180
cacctgtaaa	tttgatgggg	aatgtttaag	aattggagac	actgtgactt	gcgtctgtca	240
gttcaagtgc	aacaatgact	atgtgcctgt	gtgtggctcc	aatggggaga	gctaccagaa	300
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catgtctttg	ggctgatgtc	aagataaacac	aactacaact	actaagtctg	aagatgggca	660
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ccacatacct	tgtccggaac	attacaatgg	cttctgcctg	catgggaagt	gtgagcattc	780
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aatcgacg						908

&lt;210&gt; 351

&lt;211&gt; 472

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 351

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gatctgtcca	caacaaactt	gccctctcat	gccttgccctc	tcaccatgct	ctgctccagg	360
tcagccccct	tttggcctgt	ttgttttgtc	aaaaaccta	tctgcttctt	gcttttcttg	420
gtaatatata	tttaggggaag	atgttgcttt	gccacacac	gaagcaaagt	aa	472

&lt;210&gt; 352

&lt;211&gt; 251

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 352

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atacatggaa	aggaggggga	agccaaccca	gaaatgggct	ttctctaate	ctgggatacc	240
aataagcaca	a					251

&lt;210&gt; 353

&lt;211&gt; 436

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 353

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gtatccaaaa	gcaaaacagc	agatatata	aattaaagag	acagaagata	gacattaaca	180
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ttaacagaat	actagattca	cactggaacg	ggggtaaaga	agaaattatt	ttctataaaa	420
gggctcctaa	tgtagt					436

&lt;210&gt; 354

&lt;211&gt; 854

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 354

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&lt;210&gt; 355

&lt;211&gt; 676

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 355

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gcttaaagaa	aaccag					676

&lt;210&gt; 356

&lt;211&gt; 574

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 356

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&lt;210&gt; 357

&lt;211&gt; 393

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 357

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&lt;210&gt; 358

&lt;211&gt; 630

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 358

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&lt;210&gt; 359

&lt;211&gt; 620

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 359

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aacaaaaagc	tcacaccaaa	caaaaccatc	aacttatttt	gtattctata	acatacgaga	600
ctgtaaagat	gtgacagtgt					620

&lt;210&gt; 360

&lt;211&gt; 431

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 360

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&lt;210&gt; 361

&lt;211&gt; 351

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 361

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&lt;210&gt; 362

&lt;211&gt; 463

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 362

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cacacttgca	cacattctcc	ctgataagca	cgatgggtgtg	gacaggaagg	aaggatttca	420
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&lt;210&gt; 363

&lt;211&gt; 653

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(653)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 363

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&lt;210&gt; 364

&lt;211&gt; 401

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 364

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&lt;210&gt; 365

&lt;211&gt; 356

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 365

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&lt;210&gt; 366

&lt;211&gt; 1851

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 366

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&lt;210&gt; 367

&lt;211&gt; 668

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 367

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aaaaaaaa						668

&lt;210&gt; 368

&lt;211&gt; 1512

&lt;212&gt; DNA

&lt;213&gt; Homo sapien



&lt;400&gt; 368

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&lt;210&gt; 369

&lt;211&gt; 1853

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 369

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&lt;210&gt; 370

&lt;211&gt; 2184

&lt;212&gt; DNA

&lt;213&gt; Homo. sapien

&lt;400&gt; 370

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<210> 371  
 <211> 1855  
 <212> DNA  
 <213> Homo sapien  
  
 <220>  
 <221> misc\_feature  
 <222> (1)...(1855)  
 <223> n = A,T,C or G

<400> 371  
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&lt;210&gt; 373

&lt;211&gt; 1155

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 373

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&lt;210&gt; 374

&lt;211&gt; 2000

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 374

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 <213> Homo sapien

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&lt;210&gt; 376

&lt;211&gt; 329

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 376

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35      40      45
Leu Asp Gly Gln Gly Glu Arg Gln Glu Gln Arg Gly His Phe Trp Arg
50      55      60
Pro Gln Arg Leu Leu Cys Glu Asp Ala Trp Glu Gln Glu Val Gln Val
65      70      75      80
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85      90      95
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His Val His Gly Glu Asp Leu Asp Lys Leu His Arg Ala Ala Trp Trp
115     120     125
Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met Leu Arg Asp Thr Asp
130     135     140
Val Asn Lys Arg Asp Lys Gln Lys Arg Thr Ala Leu His Leu Ala Ser
145     150     155     160
Ala Asn Gly Asn Ser Glu Val Val Lys Leu Val Leu Asp Arg Arg Cys
165     170     175
Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr Ala Leu Thr Lys Ala
180     185     190
Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met Leu Leu Glu His Gly
195     200     205
Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn Thr Thr Leu His Tyr
210     215     220
Ala Val Tyr Asn Glu Asp Lys Leu Met Ala Lys Ala Leu Leu Leu Tyr
225     230     235     240
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245     250     255
Leu Gly Ile His Glu Gln Lys Gln Gln Val Val Lys Phe Leu Ile Lys
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Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr Gly Arg Thr Ala Leu
275     280     285
Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile Val Ser Pro Leu Leu

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 <212> PRT  
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 85 90 95  
 Glu Cys Ala Leu Met Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro  
 100 105 110  
 Asp Glu Tyr Gly Asn Thr Thr Leu His Tyr Ala Xaa Tyr Asn Glu Asp  
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 <212> PRT  
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 50 55 60  
 Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val  
 65 70 75 80  
 Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn

85 90 95  
 Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys Cys Arg Gly Ser  
 100 105 110  
 Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe  
 115 120 125  
 Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His  
 130 135 140  
 Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met  
 145 150 155 160  
 Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln Lys Arg Thr Ala  
 165 170 175  
 Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu  
 180 185 190  
 Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr  
 195 200 205  
 Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met  
 210 215 220  
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 225 230 235 240  
 Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys  
 245 250 255  
 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly  
 260 265 270  
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 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr  
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 325 330 335  
 Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val  
 340 345 350  
 Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln Met Leu Lys Ile  
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 Pro Arg Thr His Met Val Val Glu Val Asp Ser Met Pro Ala Ala Ser  
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 Ser Val Lys Lys Pro Phe Gly Leu Arg Ser Lys Met Gly Lys Trp Cys  
 405 410 415  
 Cys Arg Cys Phe Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly  
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 Thr Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys  
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 Met Gly Lys Trp Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly  
 450 455 460  
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 465 470 475 480  
 Thr Leu Arg Asn Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys  
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 Cys Arg Gly Ser Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp  
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 Asp Ser Ala Phe Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu  
 515 520 525



Asp Lys Leu His Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp  
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 545 550 555 560  
 Lys Arg Thr Ala Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val  
 565 570 575  
 Val Lys Leu Leu Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn  
 580 585 590  
 Lys Lys Arg Thr Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu  
 595 600 605  
 Cys Ala Leu Met Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp  
 610 615 620  
 Glu Tyr Gly Asn Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys  
 625 630 635 640  
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 Asn Lys His Gly Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys  
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 770 775 780  
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 Glu Glu Lys Gln Ile Glu Val Val Glu Lys Met Asn Ser Glu Leu Ser  
 995 1000 1005  
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 1315 1320 1325  
 His Glu Gln Lys Gln Gln Val Val Lys Phe Leu Ile Lys Lys Lys Ala  
 1330 1335 1340  
 Asn Leu Asn Ala Leu Asp Arg Tyr Gly Arg Thr Ala Leu Ile Leu Ala  
 1345 1350 1355 136  
 Val Cys Cys Gly Ser Ala Ser Ile Val Ser Leu Leu Leu Glu Gln Asn  
 1365 1370 1375  
 Ile Asp Val Ser Ser Gln Asp Leu Ser Gly Gln Thr Ala Arg Glu Tyr  
 1380 1385 1390  
 Ala Val Ser Ser His His His Val Ile Cys Gln Leu Leu Ser Asp Tyr  
 1395 1400 1405

Lys Glu Lys Gln Met Leu Lys Ile Ser Ser Glu Asn Ser Asn Pro Glu  
 1410 1415 1420  
 Gln Asp Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Phe Lys Gly  
 1425 1430 1435 144  
 Ser Glu Asn Ser Gln Pro Glu Lys Met Ser Gln Glu Pro Glu Ile Asn  
 1445 1450 1455  
 Lys Asp Gly Asp Arg Glu Val Glu Glu Met Lys Lys His Glu Ser  
 1460 1465 1470  
 Asn Asn Val Gly Leu Leu Glu Asn Leu Thr Asn Gly Val Thr Ala Gly  
 1475 1480 1485  
 Asn Gly Asp Asn Gly Leu Ile Pro Gln Arg Lys Ser Arg Thr Pro Glu  
 1490 1495 1500  
 Asn Gln Gln Phe Pro Asp Asn Glu Ser Glu Glu Tyr His Arg Ile Cys  
 1505 1510 1515 152  
 Glu Leu Val Ser Asp Tyr Lys Glu Lys Gln Met Pro Lys Tyr Ser Ser  
 1525 1530 1535  
 Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu Glu Glu  
 1540 1545 1550  
 Ser Gln Arg Leu Glu Gly Ser Glu Asn Gly Gln Pro Glu Lys Arg Ser  
 1555 1560 1565  
 Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp Arg Glu Leu Glu Asn Phe  
 1570 1575 1580  
 Met Ala Ile Glu Glu Met Lys Lys His Gly Ser Thr His Val Gly Phe  
 1585 1590 1595 160  
 Pro Glu Asn Leu Thr Asn Gly Ala Thr Ala Gly Asn Gly Asp Asp Gly  
 1605 1610 1615  
 Leu Ile Pro Pro Arg Lys Ser Arg Thr Pro Glu Ser Gln Gln Phe Pro  
 1620 1625 1630  
 Asp Thr Glu Asn Glu Glu Tyr His Ser Asp Glu Gln Asn Asp Thr Gln  
 1635 1640 1645  
 Lys Gln Phe Cys Glu Glu Gln Asn Thr Gly Ile Leu His Asp Glu Ile  
 1650 1655 1660  
 Leu Ile His Glu Glu Lys Gln Ile Glu Val Val Glu Lys Met Asn Ser  
 1665 1670 1675 168  
 Glu Leu Ser Leu Ser Cys Lys Lys Glu Lys Asp Ile Leu His Glu Asn  
 1685 1690 1695  
 Ser Thr Leu Arg Glu Glu Ile Ala Met Leu Arg Leu Glu Leu Asp Thr  
 1700 1705 1710  
 Met Lys His Gln Ser Gln Leu  
 1715

&lt;210&gt; 379

&lt;211&gt; 656

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 379

Met Val Val Glu Val Asp Ser Met Pro Ala Ala Ser Ser Val Lys Lys  
 1 5 10 15  
 Pro Phe Gly Leu Arg Ser Lys Met Gly Lys Trp Cys Cys Arg Cys Phe  
 20 25 30  
 Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp  
 35 40 45  
 His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Trp  
 50 55 60

Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val  
 65 70 75 80  
 Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn  
 85 90 95  
 Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys Cys Arg Gly Ser  
 100 105 110  
 Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe  
 115 120 125  
 Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His  
 130 135 140  
 Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met  
 145 150 155 160  
 Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln Lys Arg Thr Ala  
 165 170 175  
 Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu  
 180 185 190  
 Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr  
 195 200 205  
 Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met  
 210 215 220  
 Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn  
 225 230 235 240  
 Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys  
 245 250 255  
 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly  
 260 265 270  
 Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys Gln Gln Val Val  
 275 280 285  
 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr  
 290 295 300  
 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile  
 305 310 315 320  
 Val Ser Leu Leu Leu Glu Gln Asn Ile Asp Val Ser Ser Gln Asp Leu  
 325 330 335  
 Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val  
 340 345 350  
 Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln Met Leu Lys Ile  
 355 360 365  
 Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu  
 370 375 380  
 Glu Glu Ser Gln Arg Phe Lys Gly Ser Glu Asn Ser Gln Pro Glu Lys  
 385 390 395 400  
 Met Ser Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp Arg Glu Val Glu  
 405 410 415  
 Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly Leu Leu Glu Asn  
 420 425 430  
 Leu Thr Asn Gly Val Thr Ala Gly Asn Gly Asp Asn Gly Leu Ile Pro  
 435 440 445  
 Gln Arg Lys Ser Arg Thr Pro Glu Asn Gln Gln Phe Pro Asp Asn Glu  
 450 455 460  
 Ser Glu Glu Tyr His Arg Ile Cys Glu Leu Val Ser Asp Tyr Lys Glu  
 465 470 475 480  
 Lys Gln Met Pro Lys Tyr Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp  
 485 490 495  
 Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Leu Glu Gly Ser Glu

500 505 510  
 Asn Gly Gln Pro Glu Leu Glu Asn Phe Met Ala Ile Glu Glu Met Lys  
 515 520 525  
 Lys His Gly Ser Thr His Val Gly Phe Pro Glu Asn Leu Thr Asn Gly  
 530 535 540  
 Ala Thr Ala Gly Asn Gly Asp Asp Gly Leu Ile Pro Pro Arg Lys Ser  
 545 550 555 560  
 Arg Thr Pro Glu Ser Gln Gln Phe Pro Asp Thr Glu Asn Glu Glu Tyr  
 565 570 575  
 His Ser Asp Glu Gln Asn Asp Thr Gln Lys Gln Phe Cys Glu Glu Gln  
 580 585 590  
 Asn Thr Gly Ile Leu His Asp Glu Ile Leu Ile His Glu Glu Lys Gln  
 595 600 605  
 Ile Glu Val Val Glu Lys Met Asn Ser Glu Leu Ser Leu Ser Cys Lys  
 610 615 620  
 Lys Glu Lys Asp Ile Leu His Glu Asn Ser Thr Leu Arg Glu Glu Ile  
 625 630 635 640  
 Ala Met Leu Arg Leu Glu Leu Asp Thr Met Lys His Gln Ser Gln Leu  
 645 650 655

&lt;210&gt; 380

&lt;211&gt; 671

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 380

Met Val Val Glu Val Asp Ser Met Pro Ala Ala Ser Ser Val Lys Lys  
 1 5 10 15  
 Pro Phe Gly Leu Arg Ser Lys Met Gly Lys Trp Cys Cys Arg Cys Phe  
 20 25 30  
 Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp  
 35 40 45  
 His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Trp  
 50 55 60  
 Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val  
 65 70 75 80  
 Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn  
 85 90 95  
 Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys Cys Arg Gly Ser  
 100 105 110  
 Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe  
 115 120 125  
 Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His  
 130 135 140  
 Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met  
 145 150 155 160  
 Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln Lys Arg Thr Ala  
 165 170 175  
 Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu  
 180 185 190  
 Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr  
 195 200 205  
 Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met  
 210 215 220  
 Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn

225 230 235 240  
 Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys  
 245 250 255  
 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly  
 260 265 270  
 Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys Gln Gln Val Val  
 275 280 285  
 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr  
 290 295 300  
 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile  
 305 310 315 320  
 Val Ser Leu Leu Leu Glu Gln Asn Ile Asp Val Ser Ser Gln Asp Leu  
 325 330 335  
 Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val  
 340 345 350  
 Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln Met Leu Lys Ile  
 355 360 365  
 Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu  
 370 375 380  
 Glu Glu Ser Gln Arg Phe Lys Gly Ser Glu Asn Ser Gln Pro Glu Lys  
 385 390 395 400  
 Met Ser Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp Arg Glu Val Glu  
 405 410 415  
 Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly Leu Leu Glu Asn  
 420 425 430  
 Leu Thr Asn Gly Val Thr Ala Gly Asn Gly Asp Asn Gly Leu Ile Pro  
 435 440 445  
 Gln Arg Lys Ser Arg Thr Pro Glu Asn Gln Gln Phe Pro Asp Asn Glu  
 450 455 460  
 Ser Glu Glu Tyr His Arg Ile Cys Glu Leu Val Ser Asp Tyr Lys Glu  
 465 470 475 480  
 Lys Gln Met Pro Lys Tyr Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp  
 485 490 495  
 Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Leu Glu Gly Ser Glu  
 500 505 510  
 Asn Gly Gln Pro Glu Lys Arg Ser Gln Glu Pro Glu Ile Asn Lys Asp  
 515 520 525  
 Gly Asp Arg Glu Leu Glu Asn Phe Met Ala Ile Glu Glu Met Lys Lys  
 530 535 540  
 His Gly Ser Thr His Val Gly Phe Pro Glu Asn Leu Thr Asn Gly Ala  
 545 550 555 560  
 Thr Ala Gly Asn Gly Asp Asp Gly Leu Ile Pro Pro Arg Lys Ser Arg  
 565 570 575  
 Thr Pro Glu Ser Gln Gln Phe Pro Asp Thr Glu Asn Glu Glu Tyr His  
 580 585 590  
 Ser Asp Glu Gln Asn Asp Thr Gln Lys Gln Phe Cys Glu Glu Gln Asn  
 595 600 605  
 Thr Gly Ile Leu His Asp Glu Ile Leu Ile His Glu Glu Lys Gln Ile  
 610 615 620  
 Glu Val Val Glu Lys Met Asn Ser Glu Leu Ser Leu Ser Cys Lys Lys  
 625 630 635 640  
 Glu Lys Asp Ile Leu His Glu Asn Ser Thr Leu Arg Glu Glu Ile Ala  
 645 650 655  
 Met Leu Arg Leu Glu Leu Asp Thr Met Lys His Gln Ser Gln Leu  
 660 665 670

<210> 381  
 <211> 251  
 <212> DNA  
 <213> Homo sapien

<400> 381  
 ggagaagcgt ctgctggggc aggaaggggt ttccctgccc tctcaectgt ccctcaccaa 60  
 ggtaacatgc ttcccctaag ggtatcccaa cccagggggc tcaccatgac ctctgagggg 120  
 ccaatatccc aggagaagca ttggggagtt gggggcaggt gaaggacca ggactcacac 180  
 atcctggggc tccaaggcag aggagagggg cctcaagaag gtcaggagga aaatccgtaa 240  
 caagcagtca g 251

<210> 382  
 <211> 3279  
 <212> DNA  
 <213> Homo sapiens

<400> 382  
 cttectgcag ccccatgct ggtgaggggc acgggcagga acagtggacc caacatggaa 60  
 atgctggagg gtgtcaggaa gtgatcgggc tctggggcag ggaggagggg tggggagtgt 120  
 cactgggagg ggacatcctg cagaaggtag gagtgcagaa acacccgctg caggggaggg 180  
 gagagccctg cggcacctgg gggagcagag ggagcagcac ctgcccaggc ctgggaggag 240  
 gggcctggag ggcgtgagga ggagcgaggg ggctgcattg ctggagttag ggatcagggg 300  
 cagggcgcga gatggcctca cacagggaag agagggcccc tctgcaggg cctcacctgg 360  
 gccacaggag gacactgctt ttctctgag gagtgcaggag ctgtggatgg tgctggacag 420  
 aagaaggaca gggcctggc cagggtgtca gaggctgtcg ctggcttccc ttgggatca 480  
 gactgcaggg agggagggcg gcagggttgt ggggggagtg acgatgagga tgacctgggg 540  
 gtggctccag gccttgcccc tgcctggggc ctcaccagc ctccctcaca gtctcctggc 600  
 cctcagtctc tccccccac tccatcctcc atctggcctc agtgggtcat tctgatcact 660  
 gaactgacca taccagccc tgcacaggc cctccatggc tccccaatgc cctggagagg 720  
 ggacatctag tcagagagta gtcctgaaga ggtggcctct gcgatgtgcc tgtgggggca 780  
 gcacctgca gatggtcccg gccctcatcc tgetgacctg tctgcaggga ctgtcctcct 840  
 ggaccttgcc ccttgctgag gagctggacc ctgaagtccc ctccccatag gccaagactg 900  
 gagccttggt cctctgttg gactccctgc ccatattctt gtgggagtgg gttctggaga 960  
 catttctgtc tgttcctgag agctgggaat tgctctcagt catctgcctg cgcggttctg 1020  
 agagatggag ttgcctaggc agttattggg gccaatcttt ctactgtgt ctctcctcct 1080  
 ttacccttag ggtgattctg ggggtccact tgtctgtaat ggtgtgtctc aaggtatcac 1140  
 atcatggggc cctgagccat gtgcctgcc tgaagagcct gctgtgtaca ccaaggtggg 1200  
 gcattaccgg aagtggatca aggacacat cgcagccaac ccctgagtgc cctgtccca 1260  
 ccctacctc tagtaattt aagtcacat caggttctgg catcacttgg cctttctgga 1320  
 tgctggacac ctgaagctg gaactcacct ggccgaagct cgagcctcct gagtccact 1380  
 gacctgtgct ttctgggtg gagtccaggg ctgctaggaa aaggaatggg cagacacagg 1440  
 tgtatgccaa tgtttctgaa atgggtataa ttctgtcctc tccttcggaa cactggctgt 1500  
 ctctgaagac ttctcgctca gtttcagtga ggacacacac aaagacgtgg gtgacctgt 1560  
 tgtttgtggg gtgcagagat gggaggggtg gggccacccc tggaagagtg gacagtga 1620  
 caaggtggac actctctaca gatcactgag gataagctgg agccacaatg catgaggcac 1680  
 acacacagca aggttgacgc tgtaaacata gccacagctg tcctggggggc actgggaagc 1740  
 ctagataagg ccgtgagcag aaagaagggg aggtacctcc tatgtgttg aaggaggagc 1800  
 tagggggaga aactgaaagc tgattaatta caggaggttt gttcaggctc cccaaaccac 1860  
 cgtcagattt gatgatttcc tagcaggact tacagaaata aagagctatc atgctgtggg 1920  
 ttattatggg ttgttacatt gataggatac atactgaaat cagcaaaca aacagatgta 1980  
 tagattagag tgtggagaaa acagaggaaa acttgcagtt acgaagactg gcaacttggc 2040  
 ttactaagt tttcagactg gcaggaagtc aaacctatta ggctgaggac cttgtggagt 2100  
 gtagctgac cagctgatag aggaactagc cagggtggggg cctttccctt tggatggggg 2160

```

gcatatccga cagttattct ctccaagtgg agacttacgg acagcatata attctccctg 2220
caaggatgta tgataatatg tacaaagtaa ttccaactga ggaagctcac ctgataccta 2280
gtgtccaggg tttttactgg gggctctgtg gacgagtatg gactacttga ataattgacc 2340
tgaagtcctc agacctgagg ttccctagag ttcaaacaga tacagcatgg tccagagtcc 2400
cagatgtaca aaaacaggga ttcattcaca atcccatctt tagcatgaag ggtctggcat 2460
ggcccaaggc cccaagtata tcaaggcact tgggcagaac atgccaagga atcaaagtgc 2520
atctcccagg agttattcaa gggtagagcc tttacttggg atgtacaggc tttgagcagt 2580
gcagggtctg tgagtcaacc ttttattgta caggggatga gggaaaggga gaggatgagg 2640
aagccccctt ggggatttgg tttggtcttg tgatcagggt gtctatgggg ctatccctac 2700
aaagaagaat ccagaaatag gggcacattg aggaatgata ctgagcccaa agagcattca 2760
atcattgttt tatttgcctt cttttcacac cattggtgag ggagggatta ccacctggg 2820
gttatgaaga tggttgaaca cccacacat agcaccggag atatgagatc aacagtttct 2880
tagccataga gattcacagc ccagagcagg aggacgctgc acaccatgca ggatgacatg 2940
ggggatgcgc tcgggattgg tgtgaagaag caaggactgt tagaggcagg ctttatagta 3000
acaagacggt ggggcaaact ctgatttccg tgggggaatg tcatggtctt gctttactaa 3060
gttttgagac tggcaggtag tgaactcat taggctgaga accttgtgga atgcagctga 3120
cccagctgat agaggaagta gccaggtggg agcctttccc agtgggtgtg ggacatatct 3180
ggcaagattt tgtggcactc ctggttacag atactggggc agcaaataaa actgaatctt 3240
gttttcagac cttaaaaaaa aaaaaaaaaa aaaagtttt 3279

```

&lt;210&gt; 383

&lt;211&gt; 155

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 383

```

Met Ala Gly Val Arg Asp Gln Gly Gln Gly Ala Arg Trp Pro His Thr
          5                      10                      15

```

```

Gly Lys Arg Gly Pro Leu Leu Gln Gly Leu Thr Trp Ala Thr Gly Gly
          20                      25                      30

```

```

His Cys Phe Ser Ser Glu Glu Ser Gly Ala Val Asp Gly Ala Gly Gln
          35                      40                      45

```

```

Lys Lys Asp Arg Ala Trp Leu Arg Cys Pro Glu Ala Val Ala Gly Phe
          50                      55                      60

```

```

Pro Leu Gly Ser Asp Cys Arg Glu Gly Gly Arg Gln Gly Cys Gly Gly
          65                      70                      75                      80

```

```

Ser Asp Asp Glu Asp Asp Leu Gly Val Ala Pro Gly Leu Ala Pro Ala
          85                      90                      95

```

```

Trp Ala Leu Thr Gln Pro Pro Ser Gln Ser Pro Gly Pro Gln Ser Leu
          100                      105                      110

```

```

Pro Ser Thr Pro Ser Ser Ile Trp Pro Gln Trp Val Ile Leu Ile Thr
          115                      120                      125

```

```

Glu Leu Thr Ile Pro Ser Pro Ala His Gly Pro Pro Trp Leu Pro Asn
          130                      135                      140

```

```

Ala Leu Glu Arg Gly His Leu Val Arg Glu
          145                      150

```



<210> 384  
 <211> 557  
 <212> DNA  
 <213> Homo sapiens

<400> 384  
 ggatcctcta gagcgccgc ctactactac taaattcgcg gccgcgtcga cgaagaagag 60  
 aaagatgtgt ttgttttgg actctctgtg gtcccttcca atgctgtggg tttccaacca 120  
 ggggaagggg cctttttgca ttgccaagt ccataacat gagcactact ctaccatggg 180  
 tctgcctcct ggccaagcag gctggtttgc aagaatgaaa tgaatgattc tacagctagg 240  
 acttaacctt gaaatggaaa gtcttgcaat cccatttgca ggatccgtct gtgcacatgc 300  
 ctctgtagag agcagcattc ccagggacct tggaaacagt tggcactgta aggtgcttgc 360  
 tccccaaagac acatcctaaa aggtgttgta atgggtgaaaa cgtcttcctt ctttattgcc 420  
 ccttcttatt tatgtgaaca actgtttgtc tttttttgta tcttttttaa actgtaaagt 480  
 tcaattgtga aaatgaatat catgcaaata aattatgcga ttttttttcc aaagtaaaaa 540  
 aaaaaaaaaa aaaaaaa 557

<210> 385  
 <211> 337  
 <212> DNA  
 <213> Homo sapiens

<400> 385  
 ttccaggtg atgtgagagg gaagacacat ttactatcct tgatggggct gatcccttta 60  
 gtttctctag cagcagatgg gttaggagga agtgacccaa gtggttgact cctatgtgca 120  
 tctcaaagcc atctgctgtc ttcgagtacg gacacatcat cactcctgca ttgttgatca 180  
 aaacgtggag gtgcttttcc tcagctaaga agcccttagc aaaagctcga atagacttag 240  
 tatcagacag gtccagtttc cgcaccaaca cctgctggtt ccctgtcgtg gtctggatct 300  
 ctttggccac caattcccc tttccacat cccggca 337

<210> 386  
 <211> 300  
 <212> DNA  
 <213> Homo sapiens

<400> 386  
 gggcccgtc cggcccagg ccccgctcg cgagtcctcc tccccgggtg cctgcccgcga 60  
 gccgcgtcgg ccagagggg gggcggggg ctgcctctac cggctggcgg ctgtaactca 120  
 gcgaccttg cccgaaggct ctagcaagga cccaccgacc ccagccgcgg cggcggcggc 180  
 gcggactttg cccggtgtgt gggcgggagc ggactgctgt tccgcggacg ggcagcgaag 240  
 atgttagcct tcgctgccag gaccgtggac cgatcccagg gctgtggtgt aacctcagcc 300

<210> 387  
 <211> 537  
 <212> DNA  
 <213> Homo sapiens

<400> 387  
 gggccgagtc gggcaccaag ggactctttg caggcttctt tcctcggatc atcaaggctg 60  
 cccctcctg tgccatcatg atcagcacct atgagttcgg caaaagcttc ttccagaggc 120  
 tgaaccagga ccggttctg ggcggctgaa aggggcaagg aggcaaggac cccgtctctc 180  
 ccacggatgg ggagagggca ggaggagacc cagccaagt ccttttctc agcactgagg 240  
 gagggggctt gtttcccttc cctcccggcg acaagctcca gggcagggct gtccctctgg 300

```

gcgggcccagc atttcctcag acacaacttc ttctgtctgc tccagtcgtg gggatcatca 360
cttaccaccacc cccaagttc aagaccacaaat ctccagctg ccccttcgt gtttccctgt 420
gtttgctgta gctgggcatg tetccaggaa ccaagaagcc ctgagcctgg tgtagtctcc 480
ctgacccttg ttaattcctt aagtctaaag atgatgaact tcaaaaaaaaa aaaaaaa 537

```

&lt;210&gt; 388

&lt;211&gt; 520

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 388

```

aggataattt ttaaaccaat caaatgaaaa aaacaaacaa aaaaaaaagg aaatgtcatg 60
tgaggttaaa ccagtttgca ttcccctaata gtggaaaaag taagaggact actcagcact 120
gtttgaagat tgctctctct acagcttctg agaattgtgt tatttcaact gccaaagtga 180
ggacccctc cccaacatgc ccagcccac ccctaagcat ggtcccttgt caccaggcaa 240
ccaggaaact gctacttggt gacctcacca gagaccagga gggtttggt agctcacagg 300
acttcccca cccagaaga ttagcatccc atactagact cataactaac tcaactaggc 360
tcataactca ttgatgggta ttagacaatt ccatttcttt ctggttatta taaacagaaa 420
atcttctctc ttctcattac cagtaaaggc tcttggtatc tttctggttg aatgatttct 480
atgaacttgt cttattttaa tgggtgggtt ttttctggt 520

```

&lt;210&gt; 389

&lt;211&gt; 365

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 389

```

cgttgcccc gtttgacaga aggaaaggcg gagcttattc aaagtctaga gggagtggag 60
gagttaaggc tggatttcag atctgcctgg ttccagccgc agtgtgccct ctgctcccc 120
aacgacttcc caaataatct caccagcgcc ttccagctca ggcgtcctag aagcgtcttg 180
aagcctatgg ccagctgtct ttgtgttccc tctcaccgc ctgtcctcac agctgagact 240
cccaggaaac cttcagacta ccttctctg ccttcagcaa gggcggttg ccacattctc 300
tgagggtcag tggaagaacc tagactccca ttgctagagg tagaaagggg aagggtgctg 360
gggag 365

```

&lt;210&gt; 390

&lt;211&gt; 221

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(221)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 390

```

tgctctcca tctggcccc gacttctctg tcaggaaagt ggggatggac cccatctgca 60
tacacgntt ctcatgggtg tggaacatct ctgcttgccg ttccaggaag gcctctggct 120
gctctangag tctgancga ntcgttgccc cantntgaca naaggaaagg cggagcttat 180
tcaaagtcta gagggagtgg aggagttaag gctggatttc a 221

```

&lt;210&gt; 391

&lt;211&gt; 325

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(325)  
 <223> n = A,T,C or G

<400> 391  
 tggagcaggt cccgaggcct ccctagagcc tggggccgac tctgtgncga tgcangcttt 60  
 ctctcgccg cagcctggag ctgctcctgg catctaccaa caatcagncg aggcgagcag 120  
 tagccagggc actgctgcca acagccagtc cnnataccat catgtnaccc ggtgngctct 180  
 naanttingat ntccanagcc ctacccatcn tagttctgct ctcccaccgg ntaccagccc 240  
 cactgcccag gaatcctaca gccagtaccc tgtcccgcag tctctaccta ccagtacgat 300  
 gagacctccg gctactacta tgacc 325

<210> 392  
 <211> 277  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(277)  
 <223> n = A,T,C or G

<400> 392  
 atattgttta actccttctt ttatatcttt taacattttc atggngaaaag gttcacatct 60  
 agtctcactt nggcnagnn ctctacttgg agtctcttcc cgggcctggn ccagtnghaa 120  
 antaccanga accgncatgn cttaanaacn ncctgggttn tgggttnntc aatgactgca 180  
 tgcagtgcac caccctgtcc actacgtgat gctgtaggat taaagtctca cagtgggcgg 240  
 ctgaggatac agcgccgcgt cctgtgttgc tggggaa 277

<210> 393  
 <211> 566  
 <212> DNA  
 <213> Homo sapiens

<400> 393  
 actagtccag tgtggtggaa ttcgcggccg cgctcgacgga caggtcagct gtctggctca 60  
 gtgatctaca ttctgaagtt gtctgaaaat gtcttcatga ttaaattcag cctaaacggt 120  
 ttgccgggaa cactgcagag acaatgctgt gagtttccaa ccttagccca tctgcgggca 180  
 gagaaggctt agtttgtcca tcagcattat catgatatac ggactgggta cttgggttaag 240  
 gaggggtcta ggagatctgt cccttttaga gacaccttac ttataatgaa gtatttggga 300  
 ggggtgggttt caaaagtaga aatgtcctgt attccgatga tcacacctga aacattttat 360  
 catttattaa tcatccctgc ctgtgtctat tattatattc atatctctac gctggaaact 420  
 ttctgcctca atgtttactg tgcctttgtt tttgctagtt tgtgttgttg aaaaaaaaaa 480  
 cattctctgc ctgagtttta atttttgtcc aaagttattt taatctatac aattaaaagc 540  
 ttttgcttat caaaaaaaaa aaaaaa 566

<210> 394  
 <211> 384  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature

<222> (1)...(384)

<223> n = A,T,C or G

<400> 394

```
gaacatacat gtcccggcac ctgagctgca gtctgacatc atcgccatca cgggcctcgc 60
tgcaaattng gaccgggcca aggctggact gctggagcgt gtgaaggagc tacaggccna 120
gcaggaggac cgggctttaa ggagttttaa gctgagtgtc actgtagacc ccaaatacca 180
tccaagatt atcgggagaa agggggcagt aattacccaa atccggttg agcatgacgt 240
gaacatccag tttcctgata aggacgatgg gaaccagccc caggaccaa ttaccatcac 300
agggtacgaa aagaacacag aagctgccag ggatgctata ctgagaattg tgggtgaact 360
tgagcagatg gtttctgagg acgt                                     384
```

<210> 395

<211> 399

<212> DNA

<213> Homo sapiens

<400> 395

```
ggcaaaactg tgtgacctca ataagacctc gcagatccaa ggtcaagtat cagaagtgc 60
tctgaccttg gactccaaga cctacatcaa cagcctggct atattagatg atgagccagt 120
tatcagaggt ttcattcattg cggaaattgt ggagtctaag gaaatcatgg cctctgaagt 180
attcacgtct ttccagtacc ctgagttctc tatagagttg cctaacacag gcagaattgg 240
ccagctactt gtctgcaatt gtatcttcaa gaataccctg gccatccctt tgactgacgt 300
caagttctct ttggaaagcc tgggcatctc ctactacag acctctgacc atgggacggt 360
gcagcctggt gagaccatcc aatcccaaat aaaatgcac                                     399
```

<210> 396

<211> 403

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(403)

<223> n = A,T,C or G

<400> 396

```
tggagtntc agtgcaaaca agccataaag cttcagtagc aaattactgt ctcacagaaa 60
gacattttca acttctgctc cagctgctga taaaacaaat catgtgttta gcttgactcc 120
agacaaggac aacctgttcc ttcataactc tctagagaaa aaaaggagtt gttagtagat 180
actaaaaaaa gtggatgaat aatctggata ttttctctaa aaagattcct tgaaacacat 240
taggaaaatg gagggcctta tgatcagaat gctagaatta gtccattgtg ctgaagcagg 300
gtttagggga gggagtgagg gataaaagaa ggaaaaaaag aagagtgaga aaacctattt 360
atcaaagcag gtgctatcac tcaatgttag gccctgctct ttt                                     403
```

<210> 397

<211> 100

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(100)

<223> n = A,T,C or G

&lt;400&gt; 397

actagtncag tgtggtggaa ttcgcgcccg cgtcgacctt naanccatct ctatagcaaa 60  
tccatccccg ctcttggttg gtnacagaat gactgacaaa 100

&lt;210&gt; 398

&lt;211&gt; 278

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(278)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 398

gcggccgcgt cgacagcagt tccgccagcg ctgcgccctg ggtggggatg tgctgcacgc 60  
ccacctggac atctggaagt cagcggcctg gatgaaagag cggacttcac ctggggcgat 120  
tcactactgt gcttcgacca gtgaggagag ctggaccgac agcgagggtg actcatcatg 180  
ctccgggagc cccatccacc tgtggcagtt cctcaaggag ttgctactca agccccacag 240  
ctatggccgc ttcattangt ggctcaacaa ggagaagg 278

&lt;210&gt; 399

&lt;211&gt; 298

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(298)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 399

acggagggtg aggaagcgnc cctgggatcg anaggatggg tcctgncatt gaccnccctn 60  
ggggtgccng catggagcgc atgggcgcgg gcctgggcca cggcatggat cgcggtgggt 120  
ccgagatcga gcgcattggc ctggtcatgg accgcatggg ctccgtggag cgcatgggt 180  
ccggcattga gcgcattggc ccgctgggccc tcgaccacat ggccctccanc attgancgca 240  
tgggccagac catggagcgc attggctctg gcgtggagcn catgggtgcc ggcatggg 298

&lt;210&gt; 400

&lt;211&gt; 548

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 400

acatcaacta ctctctcatt ttaaggatg gcagttccct tcctccctt ttctgcctt 60  
gtacatgtac atgtatgaaa ttctctctc ttaccgaact ctctccacac atcacaagg 120  
caaagaacca cagccttaga agggtaagag ggcaccctat gaaatgaaat ggtgatttct 180  
tgagtctctt tttccacgt ttaaggggccc atggcaggac ttagagttgc gagttaagac 240  
tgcagagggc tagagaatta tttcatacag gctttgaggc caccatgtc acttatcccg 300  
tataccctct caccatcccc ttgtctactc tgatgcccc aagatgcaac tgggcagcta 360  
gttgccccca taattctggg cctttgttgt ttgttttaat tacttgggca tcccaggaag 420  
ctttccagt atctctacc atgggcccc ctctgggat caagccctc ccaggccctg 480  
tccccagccc ctctgcccc agcccccccg cttgccttgg tgctcagccc tcccattggg 540  
agcaggtt 548

<210> 401  
<211> 355  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(355)  
<223> n = A,T,C or G

<400> 401  
actgtttcca tgttatgttt ctacacattg ctacctcagt gctcctggaa acttagcttt 60  
tgatgtctcc aagtagtcca ctttcattta actctttgaa actgtatcat ctttgccaag 120  
taagagtggg ggcctatttc agctgctttg acaaaatgac tggctcctga cttaacgttc 180  
tataaatgaa tgtgctgaag caaagtgcc atggtggcgg cgaagaagan aaagatgtgt 240  
tttgttttgg actctctgtg gtcccttcca atgctgnngg ttccaacca ggggaagggt 300  
cccttttgca ttgccaagtg ccataaccat gaggactact ctaccatggn tctgc 355

<210> 402  
<211> 407  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(407)  
<223> n = A,T,C or G

<400> 402  
atggggcaag ctggataaag aaccaagacc cactggagta tgctgtcttc aagaaaccca 60  
tctcacatgc ggtggcatat ataggctcaa aataaaggaa tggagaaaaa tatttcaagc 120  
aaatggaaaa cagaaaaaag caggtgttgc actcctactt tctgacaaaa cagactatgc 180  
gaataaagat aaaaaagaga aggacattac aaaggtgggc ctgacctttg ataatctca 240  
ttgcttgata ccaacctggg ctgttttaat tgcccaaacc aaaaggataa tttgctgagg 300  
ttgtggagct tctcccttgc agagagtcct tgatctccca aaatttggtt gagatgtaag 360  
gntgattttg ctgacaactc cttttctgaa gttttactca ttccaa 407

<210> 403  
<211> 303  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(303)  
<223> n = A,T,C or G

<400> 403  
cagtatttat agccnaactg aaaagctagt agcaggcaag tctcaaattc aggcacaaaa 60  
tcctaagcaa gagccatggc atggtgaaaa tgcaaaaggga gaggctggcc aatctacaaa 120  
tagagaacaa gacctactca gtcatgaaca aaaaggcaga caccaacatg gatctcatgg 180  
gggattggat attgtaatta tagagcagga agatgacagt gatcgtcatt tggcacaaca 240  
tcttaacaac gaccgaaacc cattatttac ataaacctcc attcggtaac catgttgaaa 300  
gga 303

<210> 404  
<211> 225  
<212> DNA  
<213> Homo sapiens

<400> 404  
aagtgttaact ttttaaaaatt tagtggattt tgaaaattct tagaggaaag taaaggaaaa 60  
attgttaatg cactcattta cctttacatg gtgaaagtgc tctcttgatc ctacaaacag 120  
acattttcca ctctgtgttc catagtgtgt aagtgtatca gatgtgttg gcattgtgaat 180  
ctccaagtgc ctgtgtaata aataaagtat ctttatttca ttcat 225

<210> 405  
<211> 334  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1) ... (334)  
<223> n = A,T,C or G

<400> 405  
gagctgttat actgtgagtt ctactaggaa atcatcaaatt ctgaggggtg tctggaggac 60  
ttcaatacac ctccccccat agtgaatcag ctccaggagg gtccagtcct tctccttact 120  
tcacccccat cccatgccaa aggaagaccc tccctccttg gctcacagcc ttctctaggc 180  
ttcccagtgct ctccaggaca gagtgggtta tgttttcagc tccatccttg ctgtgagtgt 240  
ctggtgcggt tgtgcctcca gcttctgtc agtgcttcat ggacagtgtc cagcccatgt 300  
cactctccac tctctcanng tggatcccac ccct 334

<210> 406  
<211> 216  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1) ... (216)  
<223> n = A,T,C or G

<400> 406  
tttcatacct aatgagggag ttganatnac atnnaaccag gaaatgcatg gatctcaang 60  
gaaacaaaca cccaataaac tcggagtggc agactgacaa ctgtgagaca tgcacttgct 120  
acnaaacaca aatttnatgt tgcacccttg tttctacacc tgtgggttat gacaaagaca 180  
actgccaaag aatnttcaag aaggaggact gccant 216

<210> 407  
<211> 413  
<212> DNA  
<213> Homo sapiens

<400> 407  
gctgacttgc tagtatcatc tgcattcatt gaagcacaag aacttcatgc cttgactcat 60  
gtaaatgcaa taggattaaa aaataaattt gatatcacat ggaaacagac aaaaaatatt 120  
gtacaacatt gcacccagtg tcagattcta cacctggcca ctgaggaagc aagagttaat 180  
cccagaggtc tatgtcctaa tgtgttatgg caaatggatg tcatgcacgt accttcattt 240

```

ggaaaattgt catttgtcca tgtgacagtt gatacttatt cacatttcat atgggcaacc 300
tgccagacag gagaaagtct tcccatgtta aaagacattt attatcttgt ttctctgtca 360
tgggagttcc agaaaaagtt aaaacagaca atggggccagg ttctgtagta aag          413

```

&lt;210&gt; 408

&lt;211&gt; 183

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(183)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 408

```

ggagctngcc ctcaattcct ccatntctat gttancatat ttaatgtctt ttgnnattaa 60
tnccttaacta gttaatcctt aaagggctan ntaatcctta actagtcctt ccattgtgag 120
cattatcctt ccagtattcn ccttctnttt tatttactcc ttcttggtta cccatgtact 180
ntt                                     183

```

&lt;210&gt; 409

&lt;211&gt; 250

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(250)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 409

```

cccacgcacg ataagctctt tatttctgta agtcttgcta ggaaatcatc aaatctgacg 60
gtgggtttggg ggacctgaac aaacctcctg taattaatca gctttcagtt tctcccccta 120
gtccctcctt caacaacata ggaggatcct ccccttcttt ctgctcacgg ccttatctag 180
gcttcccaagt gccccagga cagcgtgggc tatgtttaca ggcntcctt gctggggggg 240
ggcctatgc                                     250

```

&lt;210&gt; 410

&lt;211&gt; 306

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(306)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 410

```

ggctgggttg caagaatgaa atgaatgatt ctacagctag gacttaacct tgaaatggaa 60
agtcttgcaa tccatttgc aggatccgtc tgtgcacatg cctctgtaga gagcagcatt 120
cccagggacc ttggaaacag ttggcactgt aagggtgctt ctccccaaga cacatcctaa 180
aagggtgttg aatggtgaaa accgcttctt tctttattgc cccttcttat ttatgtgaac 240
nactggttgg ctttttttgn atctttttta aactggaaag ttcaattgng aaaatgaata 300
tcntgc                                     306

```



<210> 411  
<211> 261  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(261)  
<223> n = A,T,C or G

<400> 411  
agagatattn cttaggtnaa agttcataga gttcccatga actatatgac tggccacaca 60  
ggatcttttg tatttaagga ttctgagatt ttgcttgagc aggattagat aaggctgttc 120  
tttaaagtgc tgaaatggaa cagatttcaa aaaaaaaccc cacaatctag ggtgggaaca 180  
aggaaggaaa gatgtgaata ggctgatggg caaaaaacca atttaccat cagttccagc 240  
cttctctcaa ggngaggcaa a 261

<210> 412  
<211> 241  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(241)  
<223> n = A,T,C or G

<400> 412  
gttcaatgtt acctgacatt tctacaacac cccactcacc gatgtattcg ttgcccagtg 60  
ggaacatacc agcctgaatt tggaaaaaat aattgtgttt cttgcccagg aaatactacg 120  
actgactttg atggctccac aaacataacc cagtgtaaaa acagaagatg tggaggggag 180  
ctgggagatt tcaactgggtta cattgaattc ccaaactacc cangcaatta ccagccaac 240  
a 241

<210> 413  
<211> 231  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(231)  
<223> n = A,T,C or G

<400> 413  
aactcttaca atccaagtga ctcactctgtg tgcttgaatc cttccactg tctcatctcc 60  
ctcatccaag tttctagtag cttctctttg ttgtgaagga taatcaaact gaacaacaaa 120  
aagtttactc tctctatttg gaacctaaaa actctcttct tctgggtct gagggctcca 180  
agaatccttg aatcanttct cagatcattg gggacaccan atcaggaacc t 231

<210> 414  
<211> 234  
<212> DNA  
<213> Homo sapiens

&lt;400&gt; 414

```
actgtccatg aagcactgag cagaagctgg aggcacaacg caccagacac tcacagcaag 60
gatggagctg aaaacataac ccactctgtc ctggaggcac tgggaagcct agagaaggct 120
gtgagccaag gagggagggt cttccttttg catgggatgg ggatgaagta aggagagggg 180
ctggaccccc tggaagctga ttcactatgg ggggaggtgt attgaagtcc tcca      234
```

&lt;210&gt; 415

&lt;211&gt; 217

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(217)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 415

```
gcataggatt aagactgagt atcttttcta cattctttta actttctaag gggcacttct 60
caaaacacag accaggtagc aaatctccac tgctctaagg ntctcaccac cactttctca 120
cacctagcaa tagtagaatt cagtcctact tctgaggcca gaagaatggt tcagaaaaat 180
antggattat aaaaaataac aattaagaaa aataatc      217
```

&lt;210&gt; 416

&lt;211&gt; 213

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(213)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 416

```
atgcataatnt aaagganact gcctcgcttt tagaagacat ctggnctgct ctctgcatga 60
ggcacagcag taaagctctt tgattcccag aatcaagaac tctccccttc agactattac 120
cgaatgcaag gtggttaatt gaaggccact aattgatgct caaatagaag gatattgact 180
atattggaac agatggagtc tctactacaa aag      213
```

&lt;210&gt; 417

&lt;211&gt; 303

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(303)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 417

```
nagtcttcag gcccatcagg gaagttcaca ctggagagaa gtcatacata tgtactgtat 60
gtgggaaagg ctttactctg agttcaaate ttcaagccca tcagagagtc cacactggag 120
agaagccata caaatgcaat gagtgtggga agagcttcag gagggattcc cattatcaag 180
ttcatctagt ggtccacaca ggagagaaac cctataaatg tgagatatgt gggaagggct 240
tcantcaaag ttcgtatctt caaatccatc ngaaggncca cagtatanan aaacctttta 300
agt      303
```

<210> 418  
<211> 328  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(328)  
<223> n = A,T,C or G

<400> 418  
tttttggcgg tgggtggggca gggacggggac angagtctca ctctgttgcc caggctggag 60  
tgcacaggca tgatctcggc tcactacaac ccctgcctcc catgtccaag cgattcttgt 120  
gcctcagcct tccctgtagc tagaattaca ggcacatgcc accacaccca gctagttttt 180  
gtatttttag tagagacagg gtttcacccat gttggccagg ctggtctcaa actcctnacc 240  
tcagnggtca ggctggtctc aaactcctga cctcaagtga tctgcccacc tcagcctccc 300  
aaagtgctan gattacaggc cgtgagcc 328

<210> 419  
<211> 389  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(389)  
<223> n = A,T,C or G

<400> 419  
cctcctcaag acggcctgtg gtccgcctcc cggcaaccaa gaagcctgca gtgccatag 60  
acccctgagc catggactgg agcctgaaag gcagcgtaca ccctgtcct gatcttgctg 120  
cttgtttcct ctctgtggct ccattcatag cacagttgtt gcactgaggc ttgtgcaggc 180  
cgagcaaggc caagctggct caaagagcaa ccagtcaact ctgccacggg gtgccaggca 240  
ccggttctcc agccaccaac ctcaactcgt cccgcaaagt gcacatcagt tcttctaccc 300  
taaaggtagg accaaagggc atctgctttt ctgaagtcct ctgctctatc agccatcacg 360  
tggcagccac tcnggctgtg tcgacgcgg 389

<210> 420  
<211> 408  
<212> DNA  
<213> Homo sapiens

<400> 420  
gttcctccta actcctgcc aaaaacagctc tcttcaacat gagagctgca cccctcctcc 60  
tggccagggc agcaagcctt agccttggtc tcttgtttct gcttttttcc tggctagacc 120  
gaagtgtact agccaaggag ttgaagtttg tgactttggt gtttcggcat ggagaccgaa 180  
gtcccattga cacctttccc actgacccca taaaggaatc ctcatggcca caaggatttg 240  
gccaactcac ccagctgggc atggagcagc attatgaact tggagagtat ataagaaaga 300  
gatatagaaa attcttgaat gagtcctata aacatgaaca ggtttatatt cgaagcacag 360  
acgttgaccg gactttgatg aagtgctatg acaaacctgg caagcccc 408

<210> 421  
<211> 352  
<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(352)

<223> n = A,T,C or G

<400> 421

```
gctcaaaaat ctttttactg atnggcatgg ctacacaatc attgactatt acggaggcca 60
gaggagaatg aggctgggcc tgggagccct gtgcctacta naagcacatt agattatcca 120
ttcactgaca gaacagggtct tttttgggtc cttcttctcc accacnatat acttgcagtc 180
ctccttcttg aagattcttt ggcagttgtc tttgtcataa cccacagggtg tagaaacaag 240
ggtgcaacat gaaatttctg tttcgtagca agtgcattgt tcacaagttg gcangtctgc 300
cactccgagt ttattgggtg tttgtttcct ttgagatcca tgcatttctt gg 352
```

<210> 422

<211> 337

<212> DNA

<213> Homo sapiens

<400> 422

```
atgccaccat gctggcaatg cagcgggcgg tcgaaggcct gcatatccag cccaagctgg 60
cgatgatcga cggcaaccgt tgcccgaagt tgccgatgcc agccgaagcg gtgggtcaagg 120
gcgatagcaa ggtgccggcg atcgccggcg cgtcaatcct ggccaagggtc agccgtgatc 180
gtgaaatggc agctgtcgaa ttgatctacc cgggttatgg catcggcggg cataagggct 240
atccgacacc ggtgcacctg gaagccttgc agcggctggg gccgacgccg attcaccgac 300
gcttcttccg ccggtacggc tggcctatga aaattat 337
```

<210> 423

<211> 310

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(310)

<223> n = A,T,C or G

<400> 423

```
gctcaaaaat ctttttactg atatggcatg gctacacaat cattgactat tagaggccag 60
aggagaatga ggcctggcct gggagccctg tgcctactan aagcncatta gattatccat 120
tactgacag aacagggtctt ttttgggtcc ttcttctcca ccacgatata cttgcagtc 180
tccttcttga agattctttg gcagttgtct ttgtcataac ccacagggtg anaaacaagg 240
gtgcaacatg aaatttctgt ttcgtagcaa gtgcatgtct cacagttgtc aagtctgccc 300
tccgagttta 310
```

<210> 424

<211> 370

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(370)

<223> n = A,T,C or G

&lt;400&gt; 424

```

gctcaaaaat ctttttactg ataggcatgg ctacacaatc attgactatt agaggccaga 60
ggagaatgag gcctggcctg ggagccctgt gcctactaga agcacattag attatccatt 120
cactgacaga acaggtcttt tttgggtcct tcttctccac cacgatatac ttgcagtcct 180
ccttcttgaa gattcttttg cagttgtctt tgtcataacc cacaggtgta gaaacatcct 240
ggttgaatct cctggaactc cctcattagg tatgaaatag catgatgcat tgcataaagt 300
cacgaagggtg gcaaagatca caacgctgcc cagganaaca ttcattgtga taagcaggac 360
tccgtcgacg

```

370

&lt;210&gt; 425

&lt;211&gt; 216

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(216)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 425

```

aattgctatn ntttattttg ccaactcaaaa taattaccaa aaaaaaaaaa tnttaaataga 60
taacaacnca acatcaagg n anaanaaca ggaatggntg acntngcata aatnggccga 120
anattatcca ttatnttaag ggttgacttc aggntacagc acacagacaa acatgcccag 180
gaggntntca ggaccgctcg atgtntntng agggagg

```

216

&lt;210&gt; 426

&lt;211&gt; 596

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 426

```

cttccagtgga ggataaccct gttgccccgg gccgagggttc tccattaggc tctgattgat 60
tggcagtcag tgatggaagg gtgttctgat cattccgact gcccgaaggg tcgctggcca 120
gctctctgtt ttgctgagtt ggcagtagga cctaatttgt taattaagag tagatgggtga 180
gctgtccttg tattttgatt aacctaattg ccttcccagc acgactcgga ttcagctgga 240
gacatcacgg caacttttaa tgaaatgatt tgaagggccca ttaagaggca cttcccgtta 300
ttaggcagtt catctgcact gataacttct tggcagctga gctggtcgga gctgtggccc 360
aaacgcacac ttggcttttg gttttgagat acaactctta atcttttagt catgcttgag 420
ggtggatggc cttttcagct ttaacccaat ttgcaactgcc ttggaagtgt agccaggaga 480
atacactcat atactcgtgg gcttagaggc cacagcagat gtcattggct tactgcctga 540
gtcccgtctg tcccatccca ggaccttcca tcggcgagta cctgggagcc cgtgct

```

596

&lt;210&gt; 427

&lt;211&gt; 107

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(107)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 427

```

gaagaattca agttagggtt attcaaaggg cttacngaga atcctanacc caggncccag 60

```

cccgggagca gccttanaga gctcctgttt gactgcccgg ctcagn

107

<210> 428

<211> 38

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(38)

<223> n = A,T,C or G

<400> 428

gaacttcna anaangactt tattcactat ttacatt

38

<210> 429

<211> 544

<212> DNA

<213> Homo sapiens

<400> 429

ctttgctgga cgggaataaaa gtggacgcaa gcatgacctc ctgatgaggg cgctgcattt 60  
attgaagagc ggctgcagcc ctgcggttca gattaaaatc cgagaattgt atagacgccg 120  
atatccacga actcttgaag gactttctga tttatccaca atcaaatacat cggttttcag 180  
tttgatggt ggctcatcac ctgtagaacc tgacttggcc gtggctggaa tccactcgtt 240  
gccttccact tcagttacac ctactcacc atcctctcct gttggttctg tgctgctca 300  
agatactaag cccacatttg agatgcagca gccatctccc ccaattctc ctgtccatcc 360  
tgatgtgcag ttaaaaaatc tgccctttta tgatgtcctt gatgttctca tcaagccac 420  
gagtttagtt caaagcagta ttcagcgatt tcaagagaag ttttttattt ttgctttgac 480  
acctcaacaa gttagagaga tatgcatatc cagggatttt ttgccagggtg gtaggagaga 540  
ttat 544

<210> 430

<211> 507

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(507)

<223> n = A,T,C or G

<400> 430

cttatcncaa tggggctccc aaacttggct gtgcagtgga aactccgggg gaattttgaa 60  
gaacactgac acccatcttc caccgccaga ctctgattta attgggctgc agtgagaaca 120  
gagcatcaat ttaaaaagct gcccagaatg ttntcctggg cagcgttggt atctttgccn 180  
ccttcgtgac tttatgcaat gcatcatgct atttcatacc taatgagga gttccaggag 240  
attcaaccag gatgtttcta cncctgtggg ttatgacaaa gacaactgcc aaagaatntt 300  
caagaaggag gactgcaagt atatcgtggt ggagaagaag gacccaaaaa agacctgttc 360  
tgtcagtga tggataatct aatgtgcttc tagtaggcac agggctccca ggccaggcct 420  
cattctctc tggcctctaa tagtcaatga ttgtgtagcc atgcctatca gtaaaaagat 480  
ttttgagcaa aaaaaaaaaa aaaaaaa 507

<210> 431

<211> 392

<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(392)  
<223> n = A,T,C or G

<400> 431  
gaaaattcag aatggataaa aacaaatgaa gtacaaaata tttcagattt acatagcgat 60  
aaacaagaaa gcacttatca ggaggactta caaatggaag tacactctan aaccatcatc 120  
tatcatggct aaatgtgaga ttagcacagc tgtattattt gtacattgca aacacctaga 180  
aagagatggg aaacaaaatc ccaggagttt tgtgtgtgga gtcctgggtt ttccaacaga 240  
catcattcca gcattctgag attagggnga ttggggatca ttctggagtt ggaatgttca 300  
acaaaagtga tggtgttagg taaaatgtac aacttctgga tctatgcaga cattgaaggt 360  
gcaatgagtc tggcttttac tctgctgttt ct 392

<210> 432  
<211> 387  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(387)  
<223> n = A,T,C or G

<400> 432  
ggtatccnta cataatcaaa tatagctgta gtacatgttt tcattggngt agattaccac 60  
aaatgcaagg caacatgtgt agatctcttg tcttattctt ttgtctataa tactgtattg 120  
ngtagtccaa gctctcggna gtccagccac tngnaaacat gtcctcttta gattaacctc 180  
gtggacnctn ttgttgnatt gtctgaactg tagngccctg tattttgctt ctgtctgnga 240  
attctgttgc ttctggggca tttccttgng atgcagagga ccaccacaca gatgacagca 300  
atctgaattg ntccaatcac agctgcgatt aagacatact gaaatcgtac aggaccggga 360  
acaacgtata gaacactgga gtccttt 387

<210> 433  
<211> 281  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(281)  
<223> n = A,T,C or G

<400> 433  
ttcaactagc anagaanact gcttcagggn gtgtaaaatg aaaggcttcc acgcagttat 60  
ctgattaaag aacactaaga gagggacaag gctagaagcc gcaggatgtc tacactatag 120  
caggcnctat ttgggttggc tggaggagct gtggaaaaca tggagagatt ggcgctggag 180  
atcgccgtgg ctattcctcn ttgntattac accagngagg ntctctgtnt gccactgggt 240  
tnnaaaaccg ntatacaata atgatagaat aggacacaca t 281

<210> 434  
<211> 484

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 434

```
ttttaaaata agcatttagt gctcagtcct tactgagtag tctttctctc ccttctcttg 60
aatttaattc tttcaacttg caatttgcaa ggattacaca tttcactgtg atgtatattg 120
tggtgcaaaa aaaaaaaagt gtctttgttt aaaattactt ggtttgtgaa tccatcttgc 180
tttttcccca ttggaactag tcattaaccc atctctgaac tggtagaaaa acatctgaag 240
agctagtcta tcagcatctg acaggtgaat tggatggttc tcagaacccat ttcaccaga 300
cagcctgttt ctatcctgtt taataaatta gtttgggttc tctacatgca taacaaaccc 360
tgctccaatc tgtcacataa aagtctgtga cttgaagttt agtcagcacc cccaccaaac 420
tttatttttc tatgtgtttt ttgcaacata tgagtgtttt gaaaataaag taccatgtc 480
ttta
```

484

&lt;210&gt; 435

&lt;211&gt; 424

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 435

```
gcgcgcgtca gaggcaggtca ctttctgcct tccagtcctt ccttcaagga agccccatgt 60
gggtagcttt caatatcgca gggtcttact cctctgcctc tataagctca aaccaccaa 120
cgatcgggca agtaaacccc ctccctcgcc gacttcggaa ctggcgagag ttcagcgcag 180
atgggcctgt ggggaggggg caagatagat gagggggagc ggcatgggtc ggggtgaccc 240
cttgagagaga ggaagaggc cacaagaggg gctgccaccg ccactaacgg agatggccct 300
ggtagagacc tttgggggtc tggaaacctt ggactcccca tgctctaact cccacactct 360
gctatcagaa acttaaactt gaggattttc tctgtttttc actcgcaata aattcagagc 420
aaac
```

424

&lt;210&gt; 436

&lt;211&gt; 667

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(667)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 436

```
accttgggaa nactctcaca atataaaggg tcgtagactt tactccaaat tccaaaaagg 60
tcctggccat gtaatcctga aagttttccc aaggtagcta taaaatcctt ataagggtgc 120
agcctcttct ggaattcctc tgatttcaaa gtctcactct caagttcttg aaaacgaggg 180
cagttcctga aaggcaggtg tagcaactga tcttcagaaa gaggaactgt gtgcaccggg 240
atgggctgcc agagtaggat aggattccag atgctgacac cttctggggg aaacaggggt 300
gccagggttg tcatagcact catcaaagtc cggtaacgt ctgtgcttcg aatataaacc 360
tgttcatgtt tataggactc attcaagaat tttctatata tctttcttat atactctcca 420
agttcataat gctgctccat gccagctgg gtgagttggc caaatccttg tggccatgag 480
gattccttta tggggtcagt gggaaagggt tcaatgggac ttgggtctcc atgccgaaac 540
accaaagtca caaacttcaa ctcttgggt agtacacttc ggtctagcca gaaaaaagg 600
agaaacaaga agccaaggct aaggcttgc gccctgccag gaggaggggt gcagctctca 660
tgttgag
```

667

&lt;210&gt; 437

&lt;211&gt; 693



&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 437

```

ctacgtctca accctcattt ttaggtaagg aatcttaagt ccaaagatat taagtgactc 60
acacagccag gtaaggaaag ctggattggc acactaggac tctaccatac cgggttttgt 120
taaagctcag gttaggaggc tgataagctt ggaaggaaact tcagacagct ttttcagatc 180
ataaaagata attcttagcc catgttcttc tccagagcag acctgaaatg acagcacagc 240
aggtactcct ctattttcac cctcttgct tctactctct ggcagtcaga cctgtgggag 300
gccatgggag aaagcagctc tctggatgtt tgtacagatc atggactatt ctctgtggac 360
catttctcca ggttacccta ggtgtcacta ttgggggggac agccagcadc tttagctttc 420
atttgagttt ctgtctgtct tcagtagagg aaacttttgc tcttcacact tcacatctga 480
acacctaact gctgttgctc ctgaggtggt gaaagacaga tatagagctt acagtattta 540
tcctatttct aggcactgag ggctgtgggg taccttgtgg tgccaaaaca gactctgttt 600
taaggacatg ttgcttcaga gatgtctgta actatctggg ggctctgttg gctctttacc 660
ctgcatcatg tgctctcttg gctgaaaatg acc

```

693

&lt;210&gt; 438

&lt;211&gt; 360

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 438

```

ctgcttatca caatgaatgt tctcctgggc agcgttgtga tctttgccac ctctgtgact 60
ttatgcaatg catcatgcta tttcatacct aatgaggagg ttccaggaga ttcaaccagg 120
atgtttctac acctgtgggt tatgacaaag acaactgcca aagaatcttc aagaaggagg 180
actgcaagta tatctggtgg agaagaagga cccaaaaaag acctgttctg tcagtgaatg 240
gataatctaa tgtgcttcta gtaggcacag ggctcccagg ccaggcctca ttctcctctg 300
gcctctaata gtcaataatt gtgtagccat gcctatcagt aaaaagattt ttgagcaaac 360

```

&lt;210&gt; 439

&lt;211&gt; 431

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(431)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 439

```

gttcctnnta actcctgcca gaaacagctc tcctcaacat gagagctgca cccctcctcc 60
tggccagggc agcaagcctt agccttgggt tcttgtttct gcttttttcc tggctagacc 120
gaagtgtact agccaaggag ttgaagtttg tgactttggt gtttcggcat ggagaccgaa 180
gtcccattga cacctttccc actgacccca taaaggaatc ctcatggcca caaggatttg 240
gccaaactcac ccagctgggc atggagcagc attatgaact tggagagtat ataagaaaga 300
gatatagaaa attcttgaat gagtctata aacatgaaca ggtttatatt cgaagcacag 360
acgttgaccg gactttgatg agtgctatga caaacctggc agcccgtcga cgcggccgcg 420
aatttagtag t

```

431

&lt;210&gt; 440

&lt;211&gt; 523

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 440

agagataaag cttaggtcaa agttcataga gttcccatga actatatgac tggccacaca 60  
ggatcttttg tattaagga ttctgagatt ttgcttgagc aggattagat aaggctgttc 120  
tttaaatgtc tgaaatggaa cagatttcaa aaaaaaaccc cacaatctag ggtgggaaca 180  
aggaaggaaa gatgtgaata ggctgatggg caaaaaacca atttaccat cagttccagc 240  
cttctctcaa ggagaggcaa agaaaggaga tacagtggag acatctggaa agttttctcc 300  
actggaaaac tgctactatc tgtttttata tttctgttaa aatatatgag gctacagaac 360  
taaaaattaa aacctctttg tgcccttggt tcttggaaca tttatgttcc ttttaaagaa 420  
acaaaaatca aactttacag aaagatttga tgtatgtaac acatatagca gctcttgaag 480  
tatatatatc atagcaaata agtcactga tgagaacaag cta 523

&lt;210&gt; 441

&lt;211&gt; 430

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 441

gttcctccta actcctgccaa gaaacagctc tcctcaacat gagagctgca cccctcctcc 60  
tggccagggc agcaagcctt agccttgaggc tcttggttct gcttttttcc tggctagacc 120  
gaagtgtact agccaaggag ttgaagtttg tgactttggt gtttcggcat ggagaccgaa 180  
gtcccattga cacttttccc actgacccca taaaggaatc ctcattggcca caaggatttg 240  
gccaactcac ccagctgggc atggagcagc attatgaact tggagagtat ataagaaaga 300  
gatatagaaa attcttgaat gagtcctata aacatgaaca ggtttatatt cgaagcacag 360  
acgttgaccg gactttgatg agtgctatga caaacctggc agcccgctga cgcggccgag 420  
aatttagtag 430

&lt;210&gt; 442

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 442

ctaaggaatt agtagtggtc ccatacattg tttggagtgt gctattctaa aagattttga 60  
tttcctggaa tgacaattat attttaactt tgggtggggga aagagttata ggaccacagt 120  
cttcacttct gatacttgta aattaatctt ttattgcact tgttttgacc attaagctat 180  
atgttttagaa atggtcattt tacggaaaaa ttagaaaaat tctgataata gtgcagaata 240  
aatgaattaa tgttttactt aatttatatt gaactgtcaa tgacaaataa aaattctttt 300  
tgattatatt ttgttttcat ttaccagaat aaaaactaag aattaaaagt ttgattacag 360  
tc 362

&lt;210&gt; 443

&lt;211&gt; 624

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(624)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 443

tttttttttt gcaacacaat atacatcaca gtgaaatgtg taatccttgc aaattgcaag 60  
ttgaaagaat taaattcaga ggagggggaga gaaagagtag tcagtaggga ctgagcacta 120  
aatgcttatt ttaaaagaaa tgtaaagagc agaaagcaat tcaggctacc ctgccttttg 180  
tgctggctag tactccggtc ggtgtcagca gcacgtggca ttgaacattg caatgtggag 240

```

cccaaaccac agaaaatggg gtgaaattgg ccaactttct attaacttgg cttcctgttt 300
tataaaatat tgtgaataat atcacctact tcaaagggca gttatgaggc ttaaataaac 360
taacgcctac aaaacactta aacatagata acataggtgc aagtactatg tatctggtac 420
atggtaaaca tccttattat taaagtcaac gctaaaatga atgtgtgtgc atatgctaata 480
agtacagaga gagggcactt aaaccaacta agggcctgga ggggaagggtt cctggaaaga 540
ngatgcttgt gctgggtcca aatcttggtc tactatgacc ttggccaaat tatttaaact 600
ttgtccctat ctgctaaaca gatc
624

```

&lt;210&gt; 444

&lt;211&gt; 425

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(425)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 444

```

gcacatcatt nntcttgcatt tctttgagaa taagaagatc agtaaatagt tcagaagtgg 60
gaagctttgt ccaggcctgt gtgtgaaccc aatgttttgc ttagaaatag aacaagtaag 120
ttcattgcta tagcataaca caaaatttgc ataagtggtg gtcagcaaat ctttgaatgc 180
tgcttaatgt gagaggttgg taaaatcctt tgtgcaacac tctaactccc tgaatgtttt 240
gctgtgctgg gacctgtgca tgccagacaa ggccaagctg gctgaaagag caaccagcca 300
cctctgcaat ctgccacctc ctgctggcag gatttgtttt tgcactcctg gaagagccaa 360
ggaggcacca gggcataagt gagtagactt atggctcgacg cggccgcgaa tttagtagta 420
gtaga
425

```

&lt;210&gt; 445

&lt;211&gt; 414

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(414)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 445

```

catgtttatg nttttggatt actttgggca cctagtgttt ctaaatcgtc tatcattctt 60
ttctgttttt caaaagcaga gatggccaga gtctcaacaa actgtatctt caagtctttg 120
tgaaattctt tgcattgtgc agattattgg atgtagtttc ctttaactag catataaatc 180
tggtgtgttt cagataaatg aacagcaaaa tgtggtggaa ttaccatttg gaacattgtg 240
aatgaaaaat tgtgtctcta gattatgtaa caaataacta tttcctaacc attgatcttt 300
ggatttttat aatcctactc acaaatagact aggttctctc tcttgtattt tgaagcagtg 360
tgggtgctgg attgataaaa aaaaaaaaaag tcgacgcggc cggaattta gtag 414

```

&lt;210&gt; 446

&lt;211&gt; 631

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(631)

<223> n = A,T,C or G

<400> 446

```
acaaattaga anaaagtgcc agagaacacc acataccttg tccggaacat tacaatggct 60
tctgcatgca tgggaagtgt gagcattcta tcaatatgca ggagccatct tgcagggtgtg 120
atgctgggta tactggacaa cactgtgaaa aaaaggacta cagtgttcta tacgttggttc 180
ccggtcctgt acgatttcag tatgtcttaa tgcagctgt gattggaaca attcagattg 240
ctgtcatctg tgtggtgggc ctctgcatca caagggccaa actttaggta atagcattgg 300
actgagattt gtaaaccttc caaccttcca ggaaatgccc cagaagcaac agaattcaca 360
gacagaagca aaatacaggg cactacagtt cagacaatac aacaagagcg tccacgaggt 420
taatctaaag ggagcatgtt tcacagtggc tggactaccg agagcttgga ctacacaata 480
cagtattata gacaaaagaa taagacaaga gatctacaca tgttgcttg catttggtgtg 540
aatctacacc aatgaaaaca tgtactacag ctatatgtga ttatgtatgg atatatttga 600
aatagtatac attgtcttga tgttttttct g                                     631
```

<210> 447

<211> 585

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(585)

<223> n = A,T,C or G

<400> 447

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cctggccatg taatcctgaa agttttccca aggtagctat aaaatcctta taagggtgca 120
gcctcttctg gaattcctct gatttcaaag tctcactctc aagttcttga aaacgagggc 180
agttcctgaa aggcaggtat agcaactgat cttcagaaag aggaactgtg tgcaccggga 240
tgggctgcca gagtaggata ggattccaga tgctgacacc ttctggggga aacagggtcg 300
ccaggtttgt catagcactc atcaaagtcc ggtcaacgtc tgtgcttga atataaacct 360
gttcatgttt ataggactca ttcaagaatt ttctatatct ctttcttata tactctccaa 420
gttcataatg ctgctccatg cccagctggg tgagttggcc aaatccttgt ggccatgagg 480
attcctttat ggggtcagtg ggaaagggtg caatgggact tgggtctcca tgccgaaaca 540
ccaaagtcac aaacttcaac tccttggtga gtacacttcg gtcta                                     585
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<210> 448

<211> 93

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(93)

<223> n = A,T,C or G

<400> 448

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ggctccctag tgccctggag agganggggc tag                                     93
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<210> 449

<211> 706

<212> DNA

<213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(706)  
 <223> n = A,T,C or G

<400> 449  
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 ttctgancac cgaactgacc atgccagccc tgccgatggc cctccatggc tccctagtgc 120  
 cctggagagg aggtgtctag tcagagagta gtccctggaag gtggcctctg ngaggagcca 180  
 cggggacagc atcctgcaga tggtcgggag cgccccattc gccattcagg ctgcgcaact 240  
 gttgggaagg gcgacgggtg cgggcctctt cgctattacg ccagctggcg aaagggggat 300  
 gtgctgcaag gcgattaagt tgggtaacgc cagggttttc ccagtcncga cgttgtaaaa 360  
 cgacggccag tgaattgaat ttaggtgacn ctatagaaga gctatgacgt cgcattgcacg 420  
 cgtacgtaag cttggatcct cttagcgggc cgcctactac tactaaattc gcggccgcgt 480  
 cgacgtggga tccncaactga gagagtggag agtgacatgt gctggacnct gtccatgaag 540  
 cactgagcag aagctggagg cacaacgcnc cagacactca cagctactca ggaggctgag 600  
 aacaggttga acctgggagg tggaggttgc aatgagctga gatcaggccn ctgcncccca 660  
 gcatggatga cagagtgaag ctccatctta aaaaaaaaaa aaaaaa 706

<210> 450  
 <211> 493  
 <212> DNA  
 <213> Homo sapiens

<400> 450  
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 aaatgaggct gagaacttta caaagggatc ttacagacat gtcgccaata tcaactgcatg 180  
 agcctaagta taagaacaac ctttggggag aaaccatcat ttgacagtga ggtacaattc 240  
 caagtcaggc agtgaaatgg gtggaattaa actcaaatta atcctgccag ctgaaacgca 300  
 agagacactg tcagagagtt aaaaagttag ttctatccat gaggtgattc cacagtcttc 360  
 tcaagtcaac acatctgtga actcacagac caagttctta aaccactgtt caaactctgc 420  
 tacacatcag aatcacctgg agagctttac aaactcccat tgccgagggt cgacgcggcc 480  
 gcgaatttag tag 493

<210> 451  
 <211> 501  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(501)  
 <223> n = A,T,C or G

<400> 451  
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 ctcttcgcta ttacgccagc tggcgaaagg gggatgtgct gcaaggcgat taagttgggt 120  
 aacgccaggg ttttcccagt cncgacgttg taaaacgacg gccagtgaat tgaatttagg 180  
 tgacnctata gaagagctat gacgtcgcat gcacgcgtac gtaagcttgg atcctctaga 240  
 gcggccgcct actactacta aattcgcgcc cgcgtcgacg tgggatccnc actgagagag 300  
 tggagagtga catgtgctgg acnctgtcca tgaagcactg agcagaagct ggaggcacia 360  
 cgcncacagc actcacagct actcaggagg ctgagaacag gttgaacctg ggagggtggag 420  
 gttgcaatga gctgagatca ggccnctgcn ccccgagcat gatgacagag tgaaactcca 480

tctttaaaaaa aaaaaaaaaa a

501

<210> 452

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(51)

<223> n = A,T,C or G

<400> 452

agacggtttc accnttataa cnccttttag gatgggnntt ggggagcaag c

51

<210> 453

<211> 317

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(317)

<223> n = A,T,C or G

<400> 453

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ttcacccana cagcctgttt ctatcctgtt taataaatta gtttgggttc tctacatgca 180  
taacaaaccc tgctccaatc tgtcacataa aagtctgtga cttgaagttt antcagcacc 240  
cccaccaaac tttatttttc tatgtgtttt ttgcaacata tgagtgtttt gaaaataagg 300  
taccatgtc tttatta 317

<210> 454

<211> 231

<212> DNA

<213> Homo sapiens

<400> 454

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taagccacgc cagctcttgc aaggagtctt gaattctcct ctgctcactc agtagaacca 120  
agaagaccaa attcttctgc atccagctt gcaaacaaaa ttgttcttct aggtctccac 180  
ccttcctttt tcagtgttcc aaagctcttc acaatttcat gaacaacagc t 231

<210> 455

<211> 231

<212> DNA

<213> Homo sapiens

<400> 455

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cattgttccg aatgggcttt ccacaggcta cacacacaaa acaggaaaca tgccaagttt 120  
gtttcaacgc attgatgact tctccaagga tcttcttttg gcatcgacca cattcagggg 180  
caaagaattt ctcatagcac agtcacaat acagggtctc tttctcctct a 231

<210> 456  
<211> 231  
<212> DNA  
<213> Homo sapiens

<400> 456  
ttggcaggta cccttacaaa gaagacacca taccttatgc gttattaggt ggaataatca 60  
ttccattcag tattatcggt attattcttg gagaaacct gtctgtttac tgtaaccttt 120  
tgcactcaaa ttcctttatc aggaataact acatagccac tatttacaaa gccattggaa 180  
cctttttatt tgggtgcagct gctagtcagt ccctgactga cattgccaag t 231

<210> 457  
<211> 231  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(231)  
<223> n = A,T,C or G

<400> 457  
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tatttgattt tattagcaat ctctttcaga agacccttga gatcattaag ctttgtatcc 180  
agttgtctaa atcgatgcct catttcctct gaggtgtcgc tggcttttgc g 231

<210> 458  
<211> 231  
<212> DNA  
<213> Homo sapiens

<400> 458  
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acaccctaac cttgggtaac agcatttgga attatcattt gggatgagta gaatttccaa 180  
ggtcctgggt taggcatttt ggggggcccag accccaggag aagaagattc t 231

<210> 459  
<211> 231  
<212> DNA  
<213> Homo sapiens

<400> 459  
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gccctgcact gttttccctc caccacagcc atcctgtccc tcattggctc tgtgctttcc 180  
actatacaca gtcaccgtcc caatgagaaa caagaaggag caccctccac a 231

<210> 460  
<211> 231  
<212> DNA  
<213> Homo sapiens

<400> 460

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cctatcaccc tattcttggg ggctgcttct tcacagtgat catgaagcct agcagcaaatt 120
cccacctccc cacacgcaca cggccagcct ggagcccaca gaagggtcct cctgcagcca 180
gtggagcttg gtccagcctc cagtccaccc ctaccaggt taaggataga a 231
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&lt;210&gt; 461

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 461

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cgaggtttga gaagctctaa tgtgcagggg agccgagaag caggcggcct agggaggggtc 60
gcgtgtgctc cagaagagtg tgtgcatgcc agaggggaaa caggcgcctg tgtgtcctgg 120
gtggggttca gtgaggagtg ggaaattggt tcagcagaac caagccgttg ggtgaataag 180
agggggattc catggcactg atagagccct atagtctcag agctgggaat t 231
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&lt;210&gt; 462

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 462

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aggtaccctc attgtagcca tgggaaaatt gatgttcagt ggggatcagt gaattaaatg 60
gggtcatgca agtataaaaa ttaaaaaaaa aagacttcat gcccattctc atatgatgtg 120
gaagaactgt tagagagacc aacagggtag tgggttagag atttccagag tcttacattt 180
tctagaggag gtatttaatt tcttctcact catccagtgt tgtatttagg a 231
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&lt;210&gt; 463

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 463

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tactccagcc tgggtgacaga gcgagaccct atcaccgccc cccacccccc caaaaaaaaa 60
actgagtaga caggtgtcct cttggcatgg taagtcttaa gtcccctccc agatctgtga 120
catttgacag gtgtcttttc ctctggacct cgggtgtccc atctgagtga gaaaaggcag 180
tggggagggt gatcttcag tcgaagcggg atagaagccc gtgtgaaaag c 231
```

&lt;210&gt; 464

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 464

```
gtactctaag attttatcta agttgccttt tctgggtggg aaagtttaac cttagtgact 60
aaggacatca catatgaaga atgtttaagt tggagggtggc aacgtgaatt gcaaacaggg 120
cctgcttcag tgactgtgtg cctgtagtcc cagctactcg ggagtctgtg tgaggccagg 180
ggtgccagcg caccagctag atgctctgta acttctaggc cccattttcc c 231
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&lt;210&gt; 465

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 465



catgttgttg tagctgtggt aatgctggct gcatctcaga caggggtaac ttcagctcct 60  
 gtggcaaatt agcaacaaat tctgacatca tatttatggg ttctgtatct ttgttgatga 120  
 aggatggcac aatttttgct tgtgttcata atatactcag attagttcag ctccatcaga 180  
 taaactggag acatgcagga cattagggta gtgttgtagc tctggtaatg a 231

<210> 466

<211> 231

<212> DNA

<213> Homo sapiens

<400> 466

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 cctgtgcaat caaatattgt ggagaattcc ctagctggag aagtcacaaa gactataggc 180  
 aataatggag accagtccca caagatgaca accagtcggt gtgtgcggt g 231

<210> 467

<211> 311

<212> DNA

<213> Homo sapiens

<400> 467

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 tgtgccttaa cagaaggctc tgagattcta agtgggaatc atttcagtga ctgtcatgtg 180  
 gcatgggtct ctgcccaagc tcgtaatgag actatagcaa ggcggctgtg ggacgtcagt 240  
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<210> 468

<211> 3112

<212> DNA

<213> Homo sapiens

<400> 468

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 tgggaaggcac tggatgcctg atgatgaagt ggactttcaa actggggcac tactgaaacg 180  
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 aaattgaact gttaacaaag gaatctctgg tctgtggtaa tggctgagca ccactgagca 660  
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&lt;210&gt; 469

&lt;211&gt; 2229

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 469

```

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aatggaatt
2229

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&lt;210&gt; 470

&lt;211&gt; 2426

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 470

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WO 00/04149 PCT/US99/15838

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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>7</sup> :</b> <b>C12N 15/12, C07K 14/47, C12Q 1/68, A61K 39/395, G01N 33/68, 33/574, C07K 16/30, C12N 15/62, 5/02 // A61P 35/00</b>		<b>A3</b>	<b>(11) International Publication Number:</b> <b>WO 00/04149</b>
<b>(21) International Application Number:</b> PCT/US99/15838		<b>(43) International Publication Date:</b> 27 January 2000 (27.01.00)	
<b>(22) International Filing Date:</b> 14 July 1999 (14.07.99)		<b>(74) Agents:</b> MAKI, David, J. et al.; Seed and Berry LLP, 6300 Columbia, 701 Fifth Avenue, Seattle, WA 98104-7092 (US).	
<b>(30) Priority Data:</b> 09/115,453 14 July 1998 (14.07.98) US 09/116,134 14 July 1998 (14.07.98) US 09/159,822 23 September 1998 (23.09.98) US 09/159,812 23 September 1998 (23.09.98) US 09/232,880 15 January 1999 (15.01.99) US 09/232,149 15 January 1999 (15.01.99) US 09/288,946 9 April 1999 (09.04.99) US		<b>(81) Designated States:</b> AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).	
<b>(71) Applicant:</b> CORIXA CORPORATION [US/US]; Suite 200, 1124 Columbia Street, Seattle, WA 98104 (US).		<b>Published</b> <i>With international search report.</i>	
<b>(72) Inventors:</b> DILLON, Davin, Clifford; 21607 N.E. 24th Street, Redmond, WA 98053 (US). HARLOCKER, Susan, Louise; 6203 20th Avenue N.W., Seattle, WA 98107 (US). YUQIU, Jiang; 5001 South 232nd Street, Kent, WA 98032 (US). XU, Jiangchun; 15805 S.E. 43rd Place, Bellevue, WA 98006 (US). MITCHAM, Jennifer, Lynn; 16677 Northeast 88th Street, Redmond, WA 98052 (US).		<b>(88) Date of publication of the international search report:</b> 20 July 2000 (20.07.00)	
<b>(54) Title:</b> COMPOSITIONS AND METHODS FOR THERAPY AND DIAGNOSIS OF PROSTATE CANCER			
<b>(57) Abstract</b>  Compositions and methods for the therapy and diagnosis of cancer, such as prostate cancer, are disclosed. Compositions may comprise one or more prostate tumor proteins, immunogenic portions thereof, or polynucleotides that encode such portions. Alternatively, a therapeutic composition may comprise an antigen presenting cell that expresses a prostate tumor protein, or a T cell that is specific for cells expressing such a protein. Such compositions may be used, for example, for the prevention and treatment of diseases such as prostate cancer. Diagnostic methods based on detecting a prostate tumor protein, or mRNA encoding such a protein, in a sample are also provided.			

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## INTERNATIONAL SEARCH REPORT

International Application No

PCT/JS 99/15838

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 C12N15/12 C07K14/47 C12Q1/68 A61K39/395 G01N33/68  
 G01N33/574 C07K16/30 C12N15/62 C12N5/02  
 //A61P35/00

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C12N C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	WO 97 33909 A (CORIXA CORP) 18 September 1997 (1997-09-18)  the whole document	1-22, 29-31, 35-49, 53-79
A	SJOGREN H O: "Therapeutic immunization against cancer antigens using genetically engineered cells" IMMUNOTECHNOLOGY, vol. 3, no. 3, 1 October 1997 (1997-10-01), pages 161-172, XP004097000 ISSN: 1380-2933 the whole document	23-28, 32-34, 53-57

☒ Further documents are listed in the continuation of box C.☒ Patent family members are listed in annex.

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\*Z\* document member of the same patent family

Date of the actual completion of the international search

31 January 2000

Date of mailing of the international search report

04.05.00

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ANDRES S.M.

## INTERNATIONAL SEARCH REPORT

International Application No

PC1, JS 99/15838

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	<p>CHU R S ET AL: "CPG OLIGODEOXYNUCLEOTIDES ACT AS ADJUVANTS THAT SWITCH ON T HELPER 1 (TH1) IMMUNITY"</p> <p>JOURNAL OF EXPERIMENTAL MEDICINE, vol. 186, no. 10, 1 November 1997 (1997-11-01), pages 1623-1631, XP002910130</p> <p>ISSN: 0022-1007</p> <p>the whole document</p> <p>---</p>	<p>14-20, 25-27, 41-47</p>
A	<p>EP 0 317 141 A (BECTON DICKINSON CO)</p> <p>24 May 1989 (1989-05-24)</p> <p>the whole document</p> <p>---</p>	<p>50-52</p>
A	<p>ZITVOGEL L ET AL: "Eradication of established murine tumors using a novel cell-free vaccine: dendritic cell-derived exosomes"</p> <p>NATURE MEDICINE, vol. 4, no. 5, 1 May 1998 (1998-05-01), pages 594-600, XP002085387</p> <p>ISSN: 1078-8956</p> <p>cited in the application</p> <p>---</p>	
P,X	<p>WO 98 37093 A (CORIXA CORP)</p> <p>27 August 1998 (1998-08-27)</p> <p>page 3, line 20 -page 22, line 2</p> <p>page 35, line 9 - last line</p> <p>page 76, line 34 -page 78, line 22</p> <p>claims</p> <p>---</p>	<p>1-15, 17-19, 21,22, 29-31, 34,35, 39-42, 44-46, 48,49, 58-79</p>
P,X	<p>WO 98 37418 A (CORIXA CORP)</p> <p>27 August 1998 (1998-08-27)</p> <p>page 2 -page 24</p> <p>example 2</p> <p>page 35, line 15 -page 36, line 11</p> <p>page 81, line 14 -page 83, line 11</p> <p>claims</p> <p>-----</p>	<p>1-15, 17-19, 21,22, 29-31, 34,35, 39-42, 44-46, 48,49, 58-79</p>

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 99/15838

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
Remark: Although claims 29-34, 48-49, 52, 55-57  
are directed to a method of treatment of the human/animal  
body, the search has been carried out and based on the alleged  
effects of the compound/composition.
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such  
an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## B x II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

see additional sheet

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all  
searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment  
of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report  
covers only those claims for which fees were paid, specifically claims Nos.:
4. ☒ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is  
restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

1-79 all partially

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

\*\*\*\*\*  
Invention 1. Claims: 1-79 (all partially)

A polypeptide comprising at least an immunogenic portion of a prostate tumor protein defined as SEQ ID 108 and which is encoded by the related SEQ IDs 2,3,107 (according to the Description of the Sequence Identifiers), fragments and variants thereof, fusion proteins comprising it, polynucleotides or oligonucleotides derived therefrom, antibodies or fragments thereof binding to the polypeptide, pharmaceutical compositions or vaccines comprising these products and their use in methods for inhibiting, monitoring or diagnosing the development of a prostate cancer, for removing tumor cells from a sample or for expanding and/or stimulating T-cells.

## Inventions 2. to 439. Claims: 1-79 (all partially and as far as applicable)

As for subject 1. but concerning respectively SEQ IDs 1,4-106,109-111,115-171,173-175,177,179-305,307-315,326,328, 330,332-335,340-375,381,382 and 384-472.

## INTERNATIONAL SEARCH REPORT

Information on patent family members

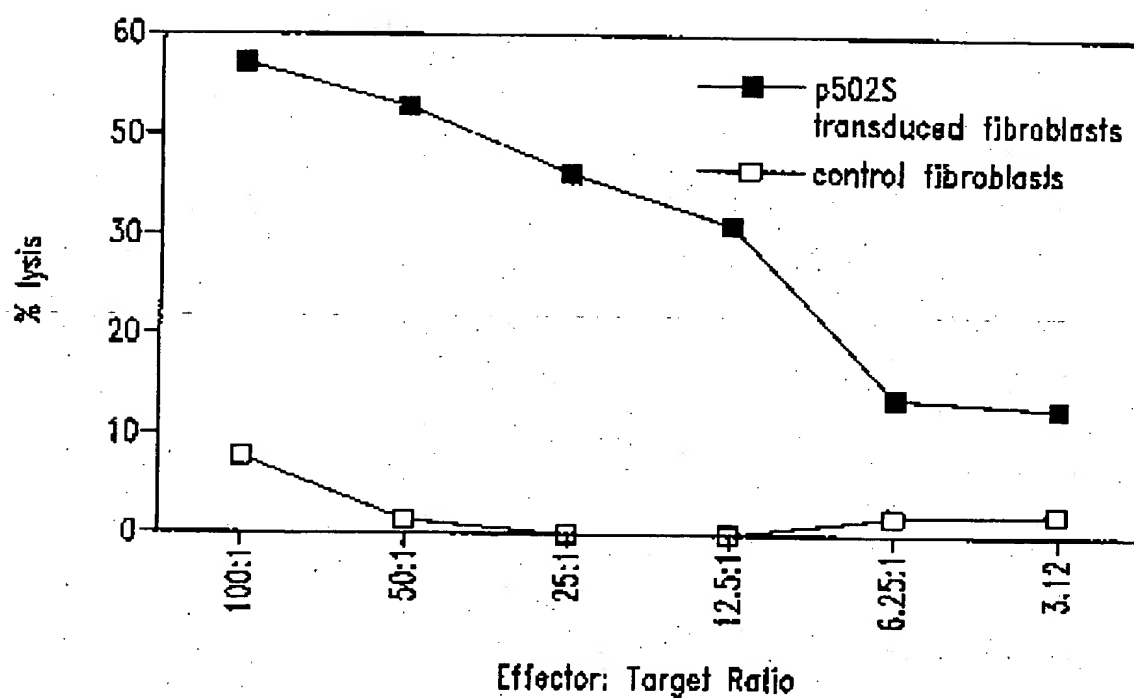
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PCT/JS 99/15838

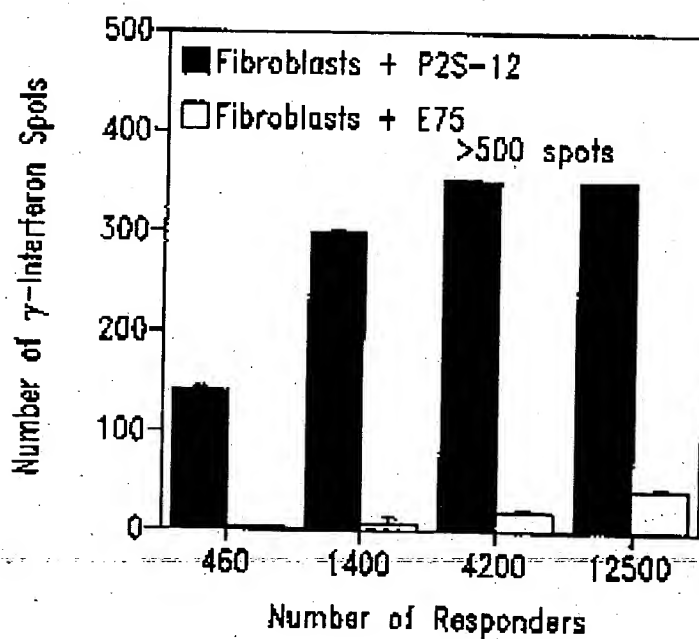
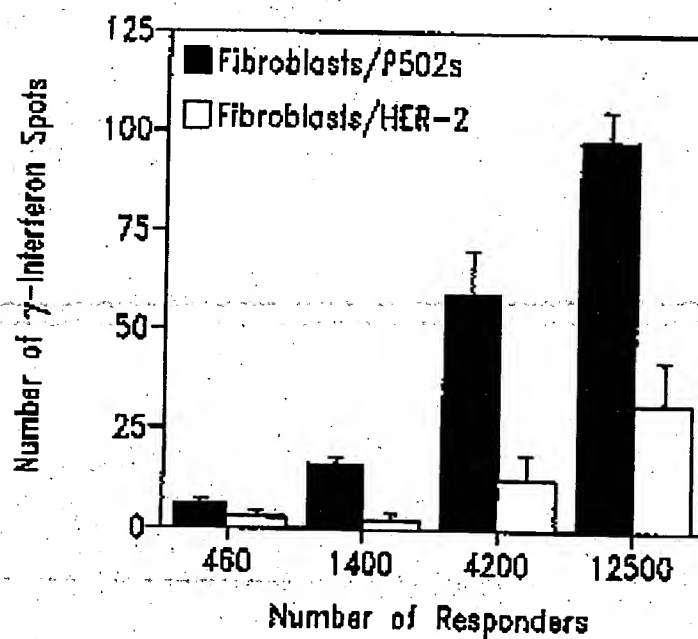
Patent document cited in search report		Publication date	Patent family member(s)		Publication date
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			BR	9708082 A	27-07-1999
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			ZA	9801536 A	08-01-1999
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*Fig. 1*

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*Fig. 2A**Fig. 2B*



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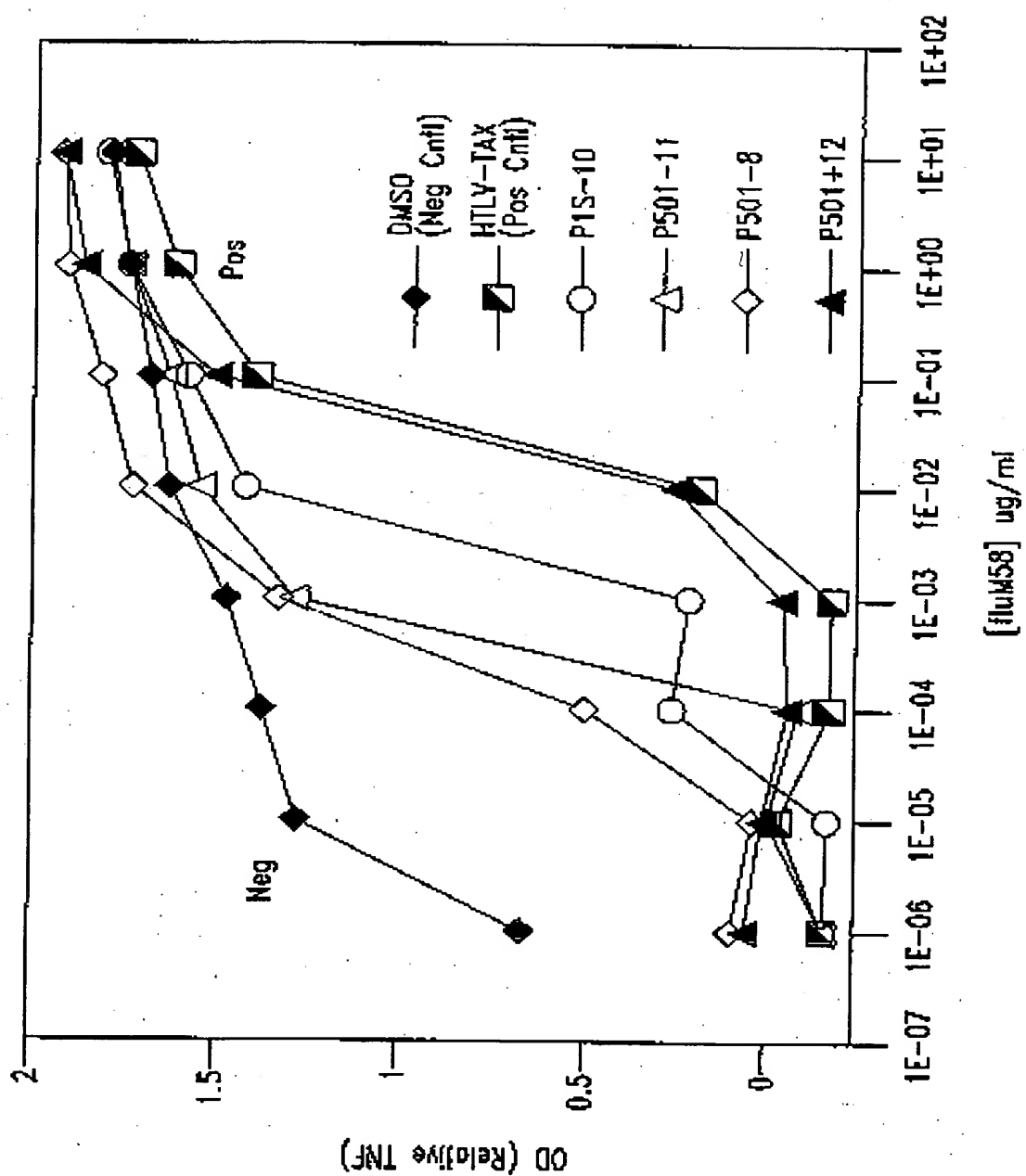
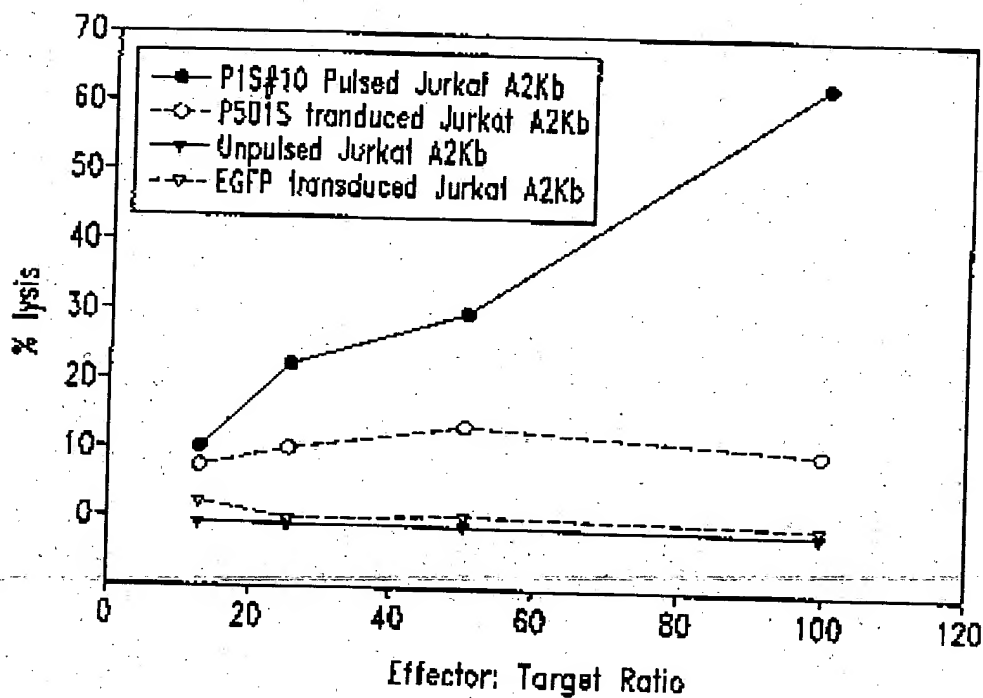
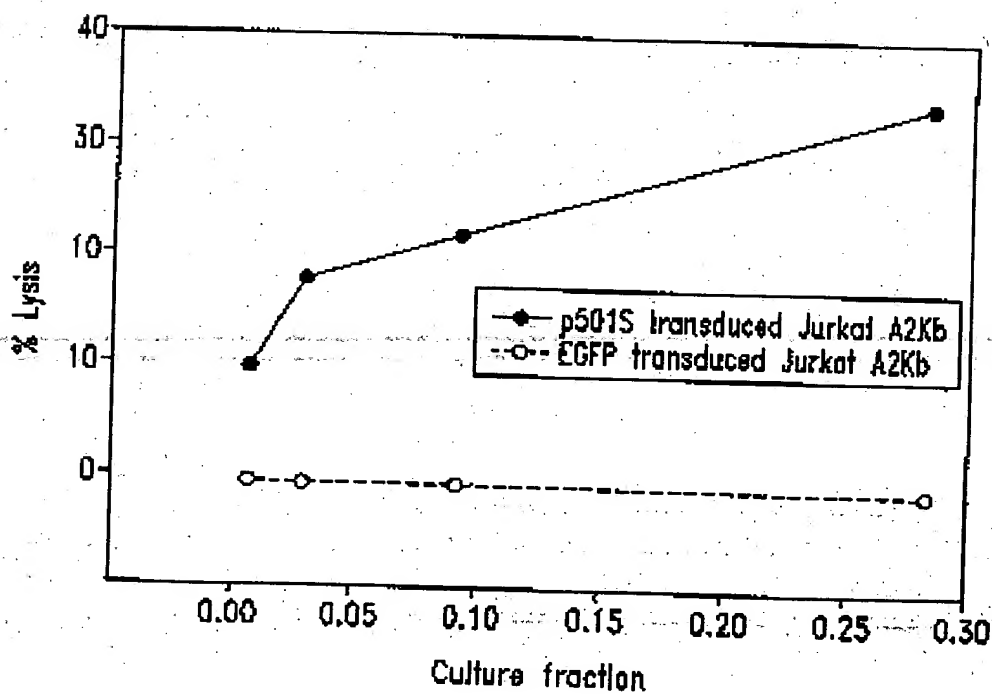


Fig. 3

4/5

*Fig. 4**Fig. 5*

5/5

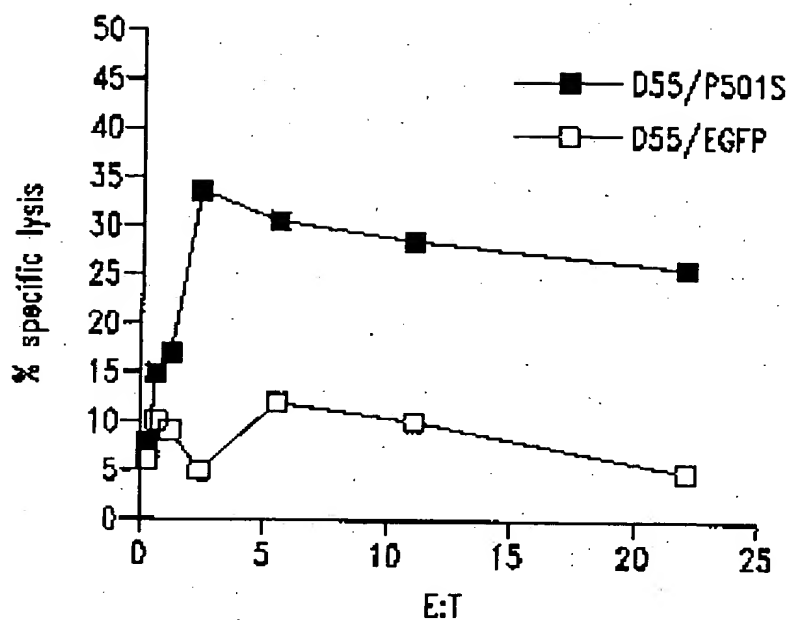


Fig. 6

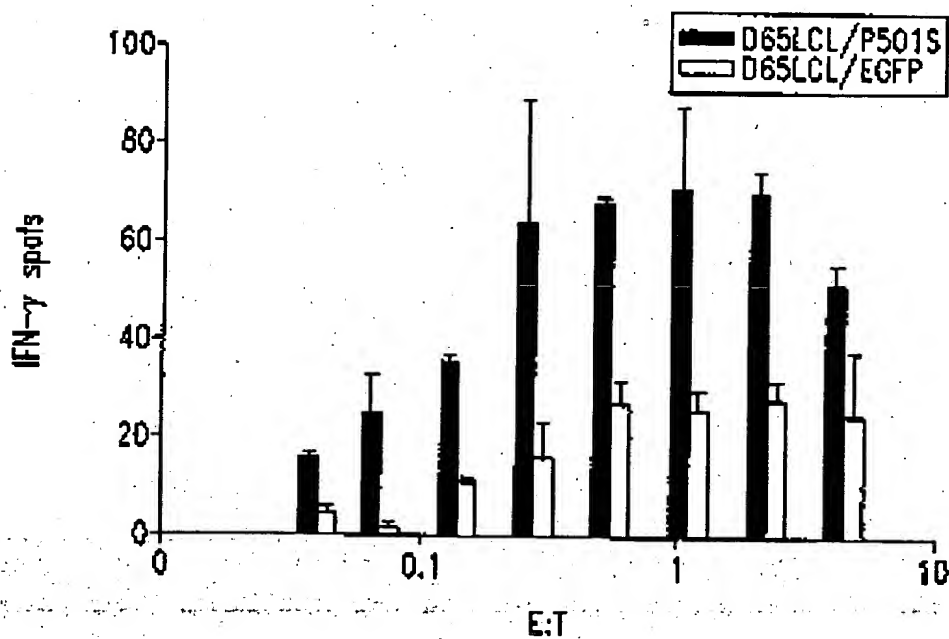


Fig. 7

SUBSTITUTE SHEET (RULE 26)

## SEQUENCE LISTING

&lt;110&gt; Corixa Corporation

<120> COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS  
OF PROSTATE CANCER AND METHODS FOR THEIR USE

&lt;130&gt; 210121.42701PC

&lt;140&gt; PCT

&lt;141&gt; 1999-07-08

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&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{816}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 2

acagaaatgt	tggatgggtg	agracctttc	tatacgactt	acaggacagc	agatggggaa	60
ttcatggctg	ttggagcaat	agaacccacg	ttctacgagc	tgtgatcaa	aggacttggg	120

ctaaagtctg	atgaacttcc	caatcagatg	aguatggatg	attggccaga	aattgaagaag	180
aagtttgcag	atgtatttgc	aaagaagacg	aaggcagagc	ggtagtcaaat	ctttgacggc	240
acagatgcct	gtgtgactcc	ggttctgaat	tttgaggagg	ttgttcatca	tgaacacaaac	300
aaggaaacggg	gctcgtttat	caccagttag	gagcaggacg	tgggcccccg	ccctgcaccc	360
ctgctgttaa	acaccccagc	catcccttct	ttcaaaaggg	atccactagt	tctaagaagcg	420
gcgcgcaccc	cggtggagct	ccagcttttg	ttcccttttag	tgggggttaa	ttgcgcgctt	480
ggcgtaatac	tggtcatagc	tgtttccctg	gtgaatttgt	tatccgctca	caattccccc	540
aacatacga	ccggaacata	aagtgttaag	cctgggggtg	ctaatgantg	agctaactcn	600
cattaattgc	gttgcgctca	ctgcgcgctt	tccagtcggg	aaaactgtcg	tgcgaatgcn	660
ttantgaatc	ngccaccccc	cgggaaaagg	cggttgcatt	ttgggcctct	tccgctttcc	720
tgcctcattg	atcctngcnc	ccggtcttcg	gctggggnga	acggttcact	ccccaagggc	780
ggctntccgg	ttatccccc	acnggggata	ccngga			816

&lt;210&gt; 3

&lt;211&gt; 773

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(773)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 3

cttttgaaag	aagggatggc	tgggggtgtt	aacagcagag	gtgcaggggcg	ggggctcacg	60
tctgtctct	cactggat	aaacgagccc	cgctccttgt	tgtgatcatg	atgaacaaac	120
tctcctaaag	tcagaacccg	agtcacacag	gcctctgtgc	cgccaaagat	ttgacaccac	180
tctgccttcg	tcttctttgc	aaatacactc	gcctccttct	tcttcatttc	tggccaatca	240
cccatgctca	tctgattggg	aagttcatca	gacttttagc	canntccttt	gattcagcagc	300
tcttagaact	ggggttctat	tgtcccaaca	gccatgaatt	ccccatctgc	tgtcctgtaa	360
gtcgtataga	aaggtgctcc	accatccaac	atgttctgtc	ctcgaggggg	ggcccgggtac	420
ccaattcgcc	ctatantgag	tctgattacg	cgcgctcact	ggcgtctgtt	ttacaaggtc	480
gtgactggga	aaacccctggg	cgttacccaa	ttatctgcct	tgcagcacat	ccccctttcg	540
ccaactgggc	gtaatanaga	aaaggcccgc	accgatcgcc	cttcccaacag	ttgcgcacct	600
gaatgggnaa	atgggacccc	cctgttaacc	cgcattnaac	ccccgcnggg	tttngttgtt	660
acccccacnt	nnaccgctta	cactttgcca	gcgccttanc	gcgcgtccc	tttnccttt	720
cttcccttcc	tttncnccn	ctttccccc	gggtttcccc	ntcaaaacc	cna	773

&lt;210&gt; 4

&lt;211&gt; 828

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(828)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 4

cctcctgaag	cctactgacc	tgtgttttct	gggtgtggagt	ccagggtctgc	taggaaaagg	60
aatgggcaga	cacaggtgtg	tgccaatgtt	totgaaatgg	gtataatttc	gtcctctcct	120
tgggaacact	ggctgtctct	gaagacttct	cgctcagttt	cagttaggac	acacacaaag	180
acgtgggtga	ccatgttgtt	tgtgggggtg	agagatggga	gggggtgggg	ccacccttga	240
agagtggaca	gtgacacaa	gtggacactc	tctacagatc	autgaggata	agrtggagcc	300
acaatgcatt	aggcacacac	acagcaaggc	tgacnotgta	aacatagccc	acgtgtcct	360

gnggggactg	ggaagcctan	atnaggccgt	gagcanaaag	aaggggagga	lccactagtt	420
ctanagcggc	cggcaccggg	gtgganctcc	ancttttgtt	cccttttagtg	agggttaatt	480
gggggcttgg	cntaatcatg	gtcatanctn	tttctgtgtg	gaaattgtta	tccgctcaca	540
attccacaca	acatacagnc	cggaaacata	aantgtaaac	clggggtgce	taatgantga	600
ctaactcaca	ttaatgtcgt	tgcgctcact	gcccgccttc	caatcnggaa	acctgtcttg	660
cnccttgcac	tnatgaatcn	gcdaadcccc	ggggaaaagc	gtttgcyttt	tgggcgctct	720
tccgcttctc	cncctantta	ntccctcnnc	tccgtccttc	cggctgcngc	aaacgggttc	780
acnctctcca	aaggggggat	tcgggtttcc	cnaatcggg	gganance		828

&lt;210&gt; 5

&lt;211&gt; 834

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (834)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 5

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attttataac	aatacaacac	tgtggctttt	aaaatttggg	tttctataga	taatttatcc	180
tgaagtaaat	ctagccatgc	ttttaaaaaa	tgttttaggt	cactccaagc	ttggcagtta	240
acatttggca	taaacastaa	taaaacaaac	acaatttaac	aaataaaca	tacaacattg	300
tagggcataa	tcatatacag	tataaggaa	aggctgtagt	gttgagtaag	cagttattag	360
aatagaatar	cttgggctct	atgcaaatat	gtctagacac	cttgattcac	tcagccctga	420
cattcagttt	tcaaaatagg	agacagggtc	tacagtatca	ttttacagtt	tccaacacat	480
tgaanaaaag	tagaaaatga	tgagttgatt	tttattaatg	cattacatcc	tcaagagtta	540
traccaaacg	ctcagttata	aaaaattttc	asgttatctt	agtcatataa	cttgggtgtgc	600
ttatttttaa	ttagtgttaa	atggattaa	tgaagacanc	aatgggtccc	taattgtgatt	660
gatattggtc	atttttlaaa	gcttctaaat	ctnaactttc	aggtttttga	actgggaacat	720
tgnatnacag	tgttccanag	tttcaacctc	ctgggaacatt	acagtggtgt	tgattcaaaa	780
tgttatcttg	ttaaaaatta	aatttttaac	tgggtggaaa	ataattttaa	atna	834

&lt;210&gt; 6

&lt;211&gt; 818

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (818)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 6

tttttttttt	tttttttttt	aagaccctca	tcaatagatg	gagacataca	gaaatagtca	60
aaccacatct	acaaaatgnc	agtatcaggu	gggggcttcg	aagccaaggt	gatgtttgga	120
tgtaaagtga	aataattagt	ggcggatgaa	gcagatagtg	aggaaagttg	agccaataat	180
gacgtgaagt	ccgtggaaag	ctgtggtctc	aaaaaatggt	gagccgtaga	tgccgtcggg	240
aatgggtgaag	ggagactcga	agtactctga	ggcttcttagg	agggtaaaaat	agagacccag	300
taaaattgta	ataagcagtg	cttgaattat	ttggtttcgg	ttgtttttta	ttagactatg	360
gtgagctcag	gtgattgata	ctcttgatgc	gagtataacg	gatgtgttta	ggagtgggac	420
ttctagggga	tttagcgggg	tgatgcctgt	tgggggcag	tgccctccta	gttggggggg	480
aggggctagg	ctggagtggt	aaaaggctca	gaaaaatcct	gcgaagaaaa	aaactttctga	540

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ggtaataaat aggattatcc cgtatcgaag gccttttttg acaggtggtg tgtggtggcc 600
ttgggtatgtg ctttctcgtg ttacatcgcg ccatacttgg tatatggtta gtgtgttggg 660
ttantanggc ctantatgaa gaacttttgg antggaatta aatcaatngc ttggccggaa 720
gtcatlanga nggctnaaaa ggccctgtta nggggtctggg ctnggtttta cccnaccat 780
ggaaatncccc ccccggaana ntgnatccct attcttaa 818

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&lt;210&gt; 7

&lt;211&gt; 817

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}... (817)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 7

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cgggccccat ttcaaaagatt tttaggggaa ttaattctag gacgatgggt atgaaactgt 120
ggtttgtctc acagatttca gagcattgac cgtagtatac ccccggtcgt gttagcgtga 180
aagtggcttg gtttagacgt ccgggaattg catctgtttt caagcctaata gtggggarag 240
ctcatgagtg caagacgtct tgtgatgtaa ttattatacn aatgggggct tcaatcggga 300
gtactactcg attgtcaacg tcaaggagtc gcaggtcgcc tggttctagg aataatgggg 360
gaagtatgta ggaattgaag attaatccgc cgtagtcggt gttctcctag gttcaatacc 420
attggtggcc aattgatttg atggtcaagg gggggtcgt tgaactcgtc tgttatgtaa 480
aggaatnctt ngygatyyga aggcnatnza ggactangga tnaatggcgg gcangatatt 540
tcaaacngtc tctanttctt gaaacgtctg aaatgttaat aenaatlaan ttngttatt 600
gaatnttngy gaaaagggct taccggacta gaaacaaat angaaanta atrntaangg 660
ctttatctnt aaaggttata accnctcta tnatcccacc caatngnatl ccccaacnna 720
acnattggat nccccanttc canaaanggc cncccccggg tgnannccnc cttttgttcc 780
cttnantgan ggttattenc cctngcntt atcanc 817

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&lt;210&gt; 8

&lt;211&gt; 799

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}... (799)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 8

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catttcgggg ttactttctt aaggaaagcc gagcgggaag tgctaactgt ggaatcggtg 60
cataaggaga actttctgct ggcacgcgct agggacagac gggagagcga ctccgagcgt 120
ctgaagcgca cgtcccagaa ggtggacttg gcactgaaac agctgggaca catccgcgag 180
tacgaacagc gcctgaaggt gctggagcgg gaggtccagc agtgtagccg cgtcctgggg 240
tgggtggcrg angcctganc cgtctgcct tgrtgcnccc angtgggccc ccacccccctg 300
acctgacctg gtccaaacac tgagccctgc tggcggactt caagganaac cccacangg 360
ggattttgct cctanantaa ggtcatctg ggcctcggcc cccccccttg gttggccttg 420
tctttgautg gagccccctg tccatctggg ccactgtcng gaccaccttt ngggagtggt 480
ctccttacaa ccacannatg ccgggtcctt cccggaaaac antcccance tnggaaggat 540
caagncctgn atccactnnt nctanaacgg gcnccnccg cngtggaaac cncctltatg 600
tcttttctnt tnaagggtta tncgccttg gccttncan ngtcctnenc ntttctnnt 660
gttnaaattg ttangcnccc nccnntccn cncnncnan cccgaccnna anntt.mann 720

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ncctgggggt nccnnncgat tgaaccnncc nccctntant tgcnttnggg nccnttgccc 780  
ctttccctct nggyanncc 799

<210> 9  
<211> 801  
<212> DNA  
<213> Homo sapien  
  
<220>  
<221> misc\_feature  
<222> (1)...(801)  
<223> n = A,T,C or G

<400> 9  
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taangatgac actcccgaag gtgggtctga cagtggcccc gatggacatg gggctccact 120  
caaggacaaag gccaccaggt gggggggucg aagcccacat gatccctact ctatgagcaa 180  
aatccccgtgt gggggcttct ccttgaagtc cggcancaggt gctcagtcct tggaccacag 240  
caggtuatgg ggttgtngnc caactggggg uoncaargca aaanggggca gggcctcngn 300  
caccataccn angacggggc taactnctg gacctccnc tccaccactt tcatgggctg 360  
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ctacatacgc ccggantcnc nctcccggtt tgtccctatc caggtncan caacaaattt 480  
cncctantg caccnattcc cacttttnc agatttcnc nccgngcttc ctntaaaag 540  
ggttgancor cggaaatnc cccaaagggt gggggccngg taaccacatn cccctnata 600  
gttgaantcc ccatnaccnn gactcnatgg anccntccnt tttaannacn tcttncactt 660  
gggaanance ctgncnctn cccctntaa tccnccctg cnangnncnt cccctnctc 720  
nccnnntng gentntnann cnaaaaaggc cennancaa tctcctnnc cctcanttcg 780  
ccancctctg aatcggrcn c 801

<210> 10  
<211> 789  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(789)  
<223> n = A,T,C or G

<400> 10  
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agatrcctgrr ctacacactg gcctccctct accaccggga gaagcaggtg ttctgccc 180  
aetaccgagg ggcactgga ggtgctagca gtgaggacag cctgatgacc agcttctgc 240  
caggccctaa gccctggagct ccttcccta atggacacgt ggggtgtgga ggcagtgccc 300  
tgctccaccr tccaccggcg ctctggggg cctctgctg tgatgtctcc gtacgtgtgg 360  
tggtgggtga gcccacgan gccagggtgg ttccggggcg gggcatctgc ctggacctcg 420  
ccatrcctgga tagtgcttcc tgetgtccca ngtggtccca tccctgttta tgggtccat 480  
tgtccagctc agccagcttg tncctgcta tatggtgtct gercagggcc tgggtctggt 540  
cccattact ttgtacaca ggtantatt gacaagaacg anttggccaa atactcagcg 600  
ttaaaaaatt ccagcaacet tgggggtgga aggcctgcct cactgggtcc aactcccg 660  
tctgttaac ccctggggc tgcggcttg gccccaatt totgttgctg ccaaanctat 720  
gtggctctct gctgccacct gttgctggct gaagtgcna cngcncant nggggggtng 780  
ggngtccc 789



<210> 11  
 <211> 772  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{772}  
 <223> n = A,T,C or G

<400> 11  
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 accaacaggc cacatcctga taagaaggtaa yaggggggtg gatcagcaaa aagaacgtgc 180  
 tctgggctga ygggacctg ttcttgtgtg ttgccccca ggaactctcc cctacaaata 240  
 actttcatac gttaaaatcc catggaggag tgtttcatcc tagaaactcc catgcaagag 300  
 ctacattaaa cgaagctgca ggttaagggg cttanagatg ggaacccagg tgactgagtt 360  
 tattcagctc ccaaaaaccc ttctctagggt gtgtctcaac taggaggcta gctgttaacc 420  
 ctgagccctgg gtaatccacc tgcagagtc cgcattcca gtccatgga cccttctggc 480  
 ctccctgtat aagtcragar tgaaccccc ttggaaggnc tccagtragg cagccctana 540  
 aactggggaa aaaaagaaag gacgccccan cccccagctg tgcancacg cactcaara 600  
 gcacagggtg gcagcaaaa aaccacttta ctctggcaca aacaaaaact ngggggggca 660  
 accccggcac ccnangggg gttaacagga oncngggnaa cntgggacc aattnaggca 720  
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<210> 12  
 <211> 751  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{751}  
 <223> n = A,T,C or G

<400> 12  
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 ttggtgtgtt tggtaggtt gtcattgcaa cagaatggg gaaaggccct gttctctttg 180  
 aagtanggtg agtccctcaa atccgtatag ttggtgaagc caccgcaatt gagcccttc 240  
 atggtggtgt tccacaattg agtgaagtc tcttggaaac cataatcttt cttagtgcca 300  
 ggcactarna gcaacgtcag ggaagtgtc agccattgtg gtgtacacca aggcgaccac 360  
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 cncggctgc gatgaagaaa tnaccccngg ttgacaaact tgcattggcac tggganccac 540  
 agtggcccca aaatcttcca aazaggatgc cccatcnatt gaccccccac atgcccactg 600  
 ccaacagggg ctgccccacn cncnnaacga tgancnatt gnacaagatc tncntggct 660  
 tnatnaact gaacctgcn tngtggctcc tgttcaggnc cngggctga cttctnaann 720  
 aahgaectcn gaagncccc ongganann g 751

<210> 13  
 <211> 729  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(729)  
 <223> n = A,T,C or G

<400> 13

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accatgcagt	gcttcagctt	cattaagacc	atgatgatcc	tcttcattt	gctnalcctt	180
ctgtgtgtg	cagccctgtt	ggcagtgggc	atctgggtgt	caatcgatga	ggcctcctt	240
ctgaagatct	tggggccact	gtcgtccagt	gccatgcagt	ttgtcaacgt	gggctacttc	300
ctcatcgag	ccggcgctgt	ggctcttagt	ctagggttcc	tgggtgtcta	tgggtgctaag	360
actgaagagc	agtgtgccc	cgtgacgttc	ttcttcctcc	tctctctcat	cttcattgct	420
gaggttgcaa	tgtgtgggtc	gccttggtgt	acaccacaat	ggctgagcac	ttcttgargt	480
tgtgtgtaat	gcttgccatc	aaaaaagat	tatgggttcc	caggaaact	tcactcaagt	540
gttggaaacac	caccatgaaa	gggtcgaagt	gctgtggtt	cnnccacta	tcaggatttt	600
gaagantcac	ctaattcaaa	gaaaaagtg	cctttccccc	attctgttg	caattgacaa	660
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attnaegg						729

<210> 14  
 <211> 816  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(816)  
 <223> n = A,T,C or G

<400> 14

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ggcaggctca	cgcagtgcc	tctgtcactg	gggaatgga	tgcgtggag	ctcgtcaaag	180
ccactcgtgt	atttttcaca	ggcagcctcg	tccgacgcct	cggggcagtt	gggggtgtct	240
tcacactcca	ggaaactgtc	natgcagcag	ccattgctgc	agcggaaactg	ggtgggtga	300
cangtgccag	agcacactgg	atggcgctt	tccatgnnan	gggcccctng	ggaaagtccc	360
tganccccc	anctgcctct	caaaagccc	accttgaca	ccccgcacag	ctagaatgga	420
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granatctgc	tccgngggg	tctantacc	anctgggaa	aagaacccc	ggngcgaac	540
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gggacaaggt	aantngccnt	ccttttaatt	cccnanctn	ccccctggtt	tggggttttn	720
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cacaaacctn	ccccaccac	gggttcngnt	ggttng			816

<210> 15  
 <211> 783  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(783)  
 <223> n = A,T,C or G

&lt;400&gt; 15

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aagaccaaa	ccaggtggaa	ctgtggggac	tcaaggaang	cactaacctg	ttccagctga	180
cagtgaclag	ctcagaccac	ccagaggara	cggccaaagt	cacagtcant	gtgctgtcca	240
ccaagcagac	agaagactac	tgcctcgcat	craacaangt	gggtcgctgc	cggggccttt	300
tcccacgctg	gtactatgac	ccracggagc	agatctgcaa	gagtttcgtt	tatggaggct	360
gcttgggcaa	caagaacaar	taccttcggg	aagaagagt	cattctancc	tgtcnggggt	420
tgcraagggtg	gcctttgana	ngcanctctg	gggtcange	gactttcccc	caggggccctt	480
ccatggaaa	grgccaacca	ntgtttctctg	gcacctgtca	gcccaccag	ttergctgca	540
ncaatggctg	ctgcatonac	antttctctg	aattgtgaca	acacccccca	ntgcccccaa	600
ccctccraac	aaagcttccc	tgttnaaaaa	tacnccantt	ggcttttnac	aaacnccccg	660
cncctccntt	ttcccncntn	acaaaagggc	ncnngenttt	gaactgccc	aaacnnggaa	720
tctnccnngg	aaaaantncc	ccccctgggt	cctnnaancc	cctucncaaa	anctncccc	780
ccc						783

&lt;210&gt; 16

&lt;211&gt; 801

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(801)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 16

gccccaatte	cagctgccac	accacccacg	gtgactgcat	tagttcggat	gtcatataaa	60
agctgattga	agcaaccctc	tactttttgg	tcgtgagcct	tttgcttggc	gcagggtttca	120
ttggctgtgt	tggtgacgtt	gtcattgcaa	cagaatgggg	gaaaggcaet	gttctctttg	180
aagtaggggtg	agtcctcaaa	atcrgtatag	ttggtgaagc	caacagcactt	gagccrtttc	240
atgggtgggtg	tcacacactg	agtgaagctt	tcctgggaac	cataatcttt	cttgetggca	300
ggcactacca	gcaacgtcag	gaagtgcaca	gccattgtgg	tgtacaccaa	ggcgaccaca	360
gcagctgcaa	cctcagcaat	gaagatgagg	aggaggatga	agaaagaacgt	cncgaggggca	420
cacttgctct	cogtcttagc	accatagcag	ccuangaac	caagagcaaa	gaccacaacg	480
cngctgrra	atgaaagaaa	ntacccacgt	tgacaaaactg	cattygccact	ggacgacagt	540
tggcccgaa	atcttcagaa	aagggatgcc	ccatcgattg	aacacccana	tgcccactgc	600
cnacagggct	gcnccnccn	gaagaatga	gccattgaag	aaggatentc	ntggtcttaa	660
tgaactgaaa	ccttgcattg	tggccctgtg	tcagggtctt	tggcagtga	ttctganaaa	720
aaggaaacngc	ntnagcccc	ccaaangana	aaacaccccc	gggtgttgcc	ctgaattggc	780
ggccaaggan	cctgcccen	g				801

&lt;210&gt; 17

&lt;211&gt; 740

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(740)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 17

gtgagagcca	ggcgtncctc	tgcctgcccc	ctcagtggca	acaccogggg	gctgttttgt	60
------------	------------	------------	------------	------------	------------	----

cctttgtgga	gctcagcag	ttccctcttt	cagaactcac	tyccnagagc	cctgaacagg	120
agccaccatg	cagtgcctca	gcttcattaa	gacctgatg	atctcttcca	atttgcctat	180
ctttctgtgt	ggtgcagccc	tgttggcagt	gggcattctg	gtgtcaatcg	atggggcctc	240
ctttctgaag	atcttcgggc	cactgtctgc	cagtgcctatg	cagtttgtca	acgtgggcta	300
cttccctcctc	gcagccggcg	ttgtggctct	tgtctcttgt	ttcctgggct	gctatgggtg	360
taagacggag	agcaagtgtg	ccctcgtgac	gttcttcttc	atccttctcc	tcactctcat	420
tgttgaaatt	gcagctgctg	tggtcgcctt	ggtgtacacc	accaatggctg	aaccttctct	480
gacgttgctg	gtantgctg	ccatcaanaa	agattatggg	ttcccaggaa	aaattcactc	540
aantntggaa	caconccatg	aaaaggggctc	caatttctgn	tggcttcccc	aactatacgg	600
gaattttgaa	agantcncct	tacttccaaa	aaaaaanant	tgccttctcc	ccctttctgc	660
tgcaatgaaa	acntcccaan	acngccaatu	aaaacctgac	cnmcaaaaa	ggntcncaaa	720
caaaaaant	maagggttn					740

&lt;210&gt; 18

&lt;211&gt; 802

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (802)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 18

ccgtcgggtg	cgctgggtcc	gngnagrcac	gaagcacgtc	agcatacaca	gctcaatca	60
caaggctctc	cagctgcgcg	acattacgca	gggcaagagc	ctccagcaac	actgcctatg	120
ggatcacatt	tactttagca	gccagggtga	caactgagag	gtgtcgaagc	ttattcttct	180
gagcctctgt	tagtggagga	agattccggg	cttcagctaa	gtgtcagcg	tatgtcccat	240
aagcaaacac	tgtgagcagc	cggaaaggtag	aggcaagtc	actctcagcc	agctctctaa	300
cattgggcat	gtccagcagt	tctcraaaca	cgtagacacc	agnggcctcc	agcactgat	360
ggatgaagtgt	ggccagcgct	gcccccttgg	ccgacttggc	tgggagcaga	aattgtctct	420
ggttctgccc	tgtcaccttc	acttergcac	tcatacactg	actgagtgtg	ggggacttgg	480
gctcaggatg	tccagagarg	tggttccguc	ccctcncctt	atgacaccgn	ccanncaacc	540
gtggctctcc	gccgntgng	tttgtctgnc	ctgggtcagg	gtctgtctgg	cncactctgc	600
aactctctgc	nggcccragg	aattcacenc	acgggaactn	gtangatcca	ctnnttctat	660
aaccggncgc	caccgcnhnt	ggaaetccac	tcttnttnc	tttacttgag	ggttaaggct	720
accctttncc	ttaccttggc	ccaaacctn	ccntgtgtcg	anattngtna	tcnggncna	780
tnccancnc	atangaagcc	ng				802

&lt;210&gt; 19

&lt;211&gt; 731

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (731)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 19

cnaagcttcc	aggtnacggg	ccgcnaence	tgaccnagg	tancanaang	cagncngcgg	60
gagccacccg	tcacngngng	gngtctttat	ngggaggggc	ggagccacat	cncaggacnt	120
cntgacccca	actcccccnc	nencantgca	gtgatgagtg	cagaaactgaa	ggtnacgttg	180
caggaaacca	gancaaannc	tgtccnntc	caagtccgcn	nagggggcgg	ggctggccac	240
gncatccnt	cnagtgtgtn	aaagccccnn	cctgtctact	tgttctggag	acngcnngga	300

catgcccagn	gttanataac	nggcngagag	tnanttttgc	tetcccttcc	ggctgcgrcn	360
ngngtntget	tagnggarat	aacctgacta	cttaactgaa	cccnngaate	tnccnccct	420
ccactaagct	cagaacaaaa	aacttcgaca	ccactcantt	gtcacttgn	tgttcaagta	480
aagtgtaccc	catncccaat	gtntgctnga	ngctctgncc	tgccttlangt	tgggtccctg	540
gaagacctat	caattnaagc	tatgtttctg	actgcctctt	gotccctgna	acaaancnacc	600
cnncnntcva	aggggggggc	ggcccccaat	ccccccaacc	ntnaatttnan	tttancccn	660
ccccnggce	cggtctttta	cnancntcn	nnacngggga	aaacnnngc	tttccccaa	720
nnatcchcc	t					731

&lt;210&gt; 20

&lt;211&gt; 754

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(754)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 20

tttttttttt	tttttttttt	taaaaacccc	ctccattnaa	tgnaaacctc	cgaaattgtc	60
caaccccctc	ntcccaatnn	ccntttccgg	gggggggttc	caaacccean	ttanotttgg	120
annttaaat	aatnttnt	tggnggnnn	anccnaatgt	nangaaagt	naaccanta	180
tnancttnaa	tncttggaac	ccngtngntt	ccaaaaatnt	ttaacctta	antccctcgg	240
aaatngttta	nygaaznccc	aantctctnt	aaggttgttt	gaaggntnaa	tnaaaanccc	300
nncccaattgt	ttttngccac	gcctgaatta	attggnttcc	gntgttttcc	nttaaaanza	360
gggnancccc	ggttantnaa	tcccccnnc	cccaattata	ccganctttt	ttngaattgg	420
gancccnccg	gaattaacgg	ggnnnttcc	tnttgggggg	cnngnncccc	ccccntccgg	480
ggttnggggc	aggnccnaat	tgtttaaggg	tccgaaaaat	ccctccnaga	aaaaaanctc	540
ccaggnctgag	nnatnggggtt	cccccccccc	canggccctt	ctcquanagt	tggggtttgg	600
ggggcctggg	attttnttcc	ccctnttccc	cccccccccc	ccnggganag	aggttngngt	660
tttngtcnnc	ggcccccncn	aaganccttt	ccgancttna	ctaaatccnt	gcctnggcga	720
agtcctttgn	agggntaaan	ggccccctnn	cggg			754

&lt;210&gt; 21

&lt;211&gt; 755

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(755)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 21

atcancccat	gaccccnacc	nnggggacnc	tcanccgggc	noncnacncc	cgggccnatca	60
ngtnagnnc	actncnnttn	natccncccc	cnccnactac	gcccncnanc	cnacgcnccta	120
nncanatncc	actganngcg	cgangtngan	ngcgaaanct	natarcanag	ncaccanacn	180
ccagctgtcc	nanaangcct	nnnatccngg	nnnatcccat	ntgnancctc	cnaggtattn	240
nnccnccan	gattttccct	anccgattac	ccntncccc	tanccctctc	cccccaacna	300
cgagggcnct	ggncnnaagg	nnccgnncnc	ccgctagntc	ccnncaagt	cnncncccta	360
aartcanccn	nattacnccg	ttcntgggta	tcactccccg	aatctcacc	tactcaactc	420
aaaaanaten	gatacaaat	aatncaagcc	tgnctatnac	actntgactg	ggctctctatt	480
ttagnggtcc	ntnaancncc	ctaatacttc	cagctctnct	tncccaattt	ccnaanggct	540
ctttcngaca	gcantttttg	gttcccnntt	gggttcttan	ngaattgccc	ttcntggaac	600

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gggtctctct tttcttctgg ttancttggg ttcttctggg cagttattat tttctctttt 660
aaattcttct ctttttcttt tggcttctta aatctctctg ctctgaaaaag gctctctggg 720
aaaaggttgt tttgaaaaa tttttgtttt gtctc 755

```

```

<210> 22
<211> 849
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(849)
<223> n = A,T,C or G

```

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<400> 22
ttttttttt tttttangtg tngtctgtga ggttagaggct tactacaant gtgaanacgt 60
acgtctngan taangcgacc cgaattctag gannctccct aatctcanac tctgaagatn 120
atctctgaaa vggaaaggtc accggnggal nttgctaggg tgnctctctc cttttctctn 180
cataacteng nggctctgac caccaccttc gggggctctg ngctctgggc cgggtctctn 240
gnnttaacct cactnngcna ccgtttctcn nctctctctg accctggcga tctgggtctn 300
tctgtctctc cctgnagcna aaaaantggg cctctggctc cttttctctc ttactaagca 360
cngctctcta nctctctgac cctctctant nnggggtgct gctctctctc cgtctctctg 420
nnaacctctn ggtctctctg gttgtctgct cctctctctg cctctctctc cnaaggaag 480
cgtctctctg gctctctctc tctctctctg nctctctctc cctctctctc cctctctctc 540
cctctctctc nctctctctc cctctctctc gctctctctc cctctctctc cctctctctc 600
nctctctctc nctctctctc cctctctctc gctctctctc cctctctctc cctctctctc 660
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nctctctctc nctctctctc cctctctctc gctctctctc cctctctctc cctctctctc 780
nctctctctc nctctctctc cctctctctc gctctctctc cctctctctc cctctctctc 840
nctctctctc nctctctctc cctctctctc gctctctctc cctctctctc cctctctctc 849

```

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<210> 23
<211> 872
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(872)
<223> n = A,T,C or G

```

```

<400> 23
ggcgaacta tactctctgc gactctctgc gctctctctc tctctctctc vgaacctg 60
tctgaanac cgtctctgac nctctctctc aagctctctc agtctctctc gactctctc 120
cactctctc agctctctc nctctctctc aagctctctc attctctctc agctctctc 180
nggctctctc tactctctc nctctctctc aagctctctc gttctctctc cgtctctctc 240
ctctctctc tactctctc nctctctctc aagctctctc gttctctctc cgtctctctc 300
tctctctctc nctctctctc cctctctctc cctctctctc nctctctctc cgtctctctc 360
nctctctctc nctctctctc cctctctctc cctctctctc nctctctctc cgtctctctc 420
accctctctc cctctctctc cctctctctc cctctctctc nctctctctc cgtctctctc 480
tctctctctc tctctctctc gttctctctc nctctctctc cctctctctc cgtctctctc 540
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cgtctctctc cgtctctctc nctctctctc nctctctctc cctctctctc cgtctctctc 660
cctctctctc gctctctctc ggtctctctc ggtctctctc nctctctctc cgtctctctc 720
cgtctctctc cgtctctctc cgtctctctc cgtctctctc cgtctctctc cgtctctctc 780

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ntctcccccg ngngcncntc tcaguctonc cccccccct ctctycantg tntctctctc 840  
tnaccnntac gantnttcgn cccctctctt cc 872

<210> 24  
<211> 815  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(815)  
<223> n = A,T,C or G

<400> 24  
gcattgcaagc ttgagtactc tatagngton cctaatanc ttggontaat catgggtona 60  
nctgncctcc tgtgtcaaat gtatacnaaa tanatatgaa tctnatnba caaganngtc 120  
tcntncattt gtaacaantg tontgtccat cctgtcngan canatterca tnnattncgn 180  
cgcattcncn gncantatn taatngggaa ntcnnntnon ncccnncat ctatctncc 240  
gcnccctgac tggagagat ggatnattc tntntgacc nacatgttc tcttggattn 300  
aanaccccc cgcngncac cggttngng cnagccnntc ccaagaccc ctgtggaggt 360  
aacctgggtc agannccatc aacntgggaa acccgcnnc angbnnagt ngnnncan 420  
gatercgtrc agnntnacc atcccttcc agcgccccct ttngtgccct anagngnagc 480  
gtgtccnanc vncatcaat ganacgcgc agnccancc caatlnggca caatgtcngc 540  
gaacccccca gggggantna tncaanccr caggattgtc vncncngaa atcccnccnc 600  
ccnccctac cccncttgg gacngtgacc aanteccgga gtnccagtc ggcngnctc 660  
cccccccggt nncntgggg ggggtgaact cngnntcanc cngncgaggn ntcgnaagga 720  
accgncctn ggcggaanng ancnntcnga agngccnnt cgtatacccc cccctcncca 780  
nccnccngnt agntccccc cngggtnccg aangg 815

<210> 25  
<211> 775  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> (1)...(775)  
<223> n = A,T,C or G

<400> 25  
ccgagatgtc tcgctccgta gacttagctg tgcctcgact actctctctt tctggcctgg 60  
aggctatcca ycgtaactca agattccgg tttactcacg tcactccaga gagaatggaa 120  
agtaaatctt cctgaattgc tatgtgtctg ggtttcatcc atccgacatt gaenttgact 180  
tactgaagaa tgganagaga attgaaaaay tggagcattc agacttgtct ttcagcaagg 240  
actggctctt ctatctctg tactacactg aattccccc tactgaaaa gatgagtatg 300  
cctgacctgt gaaccatgtg actttgtcac agcccaagat agttaagtgg gatcgagaca 360  
tgtaaccaan cncatggaa gtttgaagat ggcgcatttg gattgagatga attccaatl 420  
ctgcttgcct gcntttlaat antgatatgc ntatccccc taccctttat gncccraat 480  
tgtagggttc acatnangt tcnctngga catgatctc ctttataant cncncttcy 540  
aattgcccgt cncnngttn ngaatgttcc cnaaccacg gttggctccc ccaggctncc 600  
tcttaggaa gggcctgggc cnccttncan ggttggggga accnaaatc tcncttntgc 660  
cncnccccc cmtcttngg nncnctt tggaaacctc cnatccccc tggctcnna 720  
ncttuncta anaaacttn aaanogtngc naaanntttn acttccccc ttacc 775

<210> 26

&lt;211&gt; 820

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(820)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 26

anattantac	agtgtaatct	tttcccaagag	gtgtgtanag	ggaacggggc	ctagaggcat	60
cccaagata	nottatanca	acagtgtttt	gaccaaagag	tgctgggcac	atttctgtca	120
gaaaaggtgg	cggctcccat	cactctctct	ctcccatagc	catccagag	gggtgagtag	180
ccatcangcc	ttcggtggga	gggagtcang	gaacaaacaa	accacagagc	anacagacca	240
ntgatgacca	tgggcggggag	cgagctctct	ccctgnacrg	gggtggcana	nganagccta	300
netgaggggt	cacactataa	acgttaacga	cmagatnan	caactgtctc	aagtgcaccc	360
ttctacactg	acnaccagng	acnnnaact	gengcctggg	garagcncctg	ggancagcta	420
acnnagcact	cactgtcccc	cccatggccg	tnccntccc	tygtccctgnc	aaggggaagct	480
ccctgttggg	attncgggga	naccaaagga	ccccctctct	ccancgtgga	aggaaaaann	540
gatggaattt	tncccttcgg	gcctntccc	tcttcttta	cagcccccct	ntactctctc	600
tcctctctct	ntctctnenc	acttttnacc	ccnnnatttc	ccttnatttg	tcggannctn	660
ganattccac	tnnccgctnc	ctnctatcng	naaacnaaa	nactntctna	ccnnggggat	720
gggnccctcg	ntcatctctc	cttttctnct	acnccnctt	ctttgcctct	cttngatcaa	
7B0tccaaacntc	gntggcctn	ccccccnnn	tccttttccc			

820

&lt;210&gt; 27

&lt;211&gt; 818

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(818)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 27

tctgggtgat	ggctctctcc	tcctcagggg	cctctgactg	ctctgggcca	aagaatctct	60
tgcttctctc	cagagcccca	ggcagcgggt	attcagccct	gcccacactg	attctgatga	120
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ctgctgagca	cttccgcccc	tcacccctgc	cagcccttgc	catagctctc	gggctgggtc	240
tcacccctca	gggttctgct	cttccangcc	ngccanccag	tggtgctggg	ccacactggc	300
ttcttctctc	cccttccctg	gotctganc	tctgtctctc	tgtctgtgct	angcnccttg	360
gctctcagtt	tccttctctc	anngaactct	gtttctgann	tcttctctta	actntgantt	420
tatnaccnan	tggctgtctc	tgctnacttc	taatgggccc	gacccgctaa	tccttccctc	480
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ctcccttctc	ctnaccangg	gcnnnnaccg	ccctnctctn	ggggggcang	gttctctnnc	600
ctgctnacc	cctctcnnnt	tnctctgtct	cnnccnccgc	angcannctc	ncngtcccn	660
tnctctctcn	ngntctgnaa	ngntctnctn	tnnnnnngcn	ngntnctn	tcctctctnc	720
cnnctgnaag	tnntttnnnc	ncngnncccc	nnnnccnnnn	ngnnctn	tcctctctnc	780
ccnnccccc	ngnattcagg	cctccnctct	cgggcctc			818

&lt;210&gt; 28

&lt;211&gt; 731

&lt;212&gt; DNA



<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(731)

<223> n = A,T,C or G

<400> 28

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tcccaacatg	anggtgnngt	tctcttttga	angaggggtg	ngtttttann	ccnggtgggt	120
gattnaaccc	catgtgatgg	agnnaaagg	tttnagggat	ttttcggctc	ttatcagtat	180
ntanattcct	gtnaatcgga	aatnatntt	tannngyaa	aatnttgctc	ccatccgnaa	240
atttctcccg	ggtagtgcac	nttngggggn	cngccangtt	tcccaggctg	ctanaatngt	300
actaaagntt	naagtgggan	tncaaatgaa	aacctnnuac	agagnatccn	taccgactg	360
tnntttncct	tggccctntg	actctgcnn	agcccaatar	ccnngngnat	gtcnccmgn	420
nnngcggnnc	tgaannnnnc	tgnnggeton	gancatcang	gggtttcgca	tcaaaagcnn	480
cgtttencat	naaggcaatt	tngcctcacc	caaccnctng	ccctcnncca	tttngccgtc	540
nggttctnct	acgtctntng	cnctctnnnt	ganattttnc	ccgcttnggg	naancctcct	600
gnaatgggta	gggncttntc	ttttnaccnn	gnggtntact	aatcnnctnc	acgtctnctt	660
tctcnacccc	cccccttttt	caatccacnc	ggcnaatggg	gtctccccnn	cgangggggg	720
nnnccannnc	c					731

<210> 29

<211> 822

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(822)

<223> n = A,T,C or G

<400> 29

actagtccag	tgtgggtggaa	ttccattgtg	ttgggggnnc	ttctatgant	antnttagat	60
cgctcanacc	tccanccctc	ccnaccnangc	ctataangaa	naanaataga	netgtcnntt	120
atntntacnc	tcatanncct	cnnaaccacc	tccctcttaa	ccctactgtt	gcctatngcn	180
tnnctantct	ntggcgccctn	cnanccaccn	gtggggcnac	cnongynatt	ctcnatctcc	240
tenccatntn	gcctananta	ngtncatccc	ctataccctac	cccaatgcta	nnnctaaanc	300
tccatnantt	annataacta	ccactgaent	ngactttcnc	atnanctcct	aatttgaaac	360
tactctgact	cccacngcct	annattagc	ancttccccc	nacnatntct	caacczaate	420
ntcaaccaac	tactctantg	ttcnccaacc	nttncctccg	atcccccnaa	aacccccctc	480
ccaaataccc	nccacctgac	ncctaaccnn	caccatcccg	gcaagccnnc	gynccatttan	540
ccactggagt	cacnatngga	naaaaaaac	cnnaactctc	tancnennat	ctccctaaana	600
aatnctcctn	naatttactn	ncantnccat	caanccacn	tgaaccnnaa	ccccgttttt	660
tanatccctt	ctttcgaaaa	ccnacccttt	annncccaac	ctttnggggc	ccccnctnc	720
ccnaatgaag	gnccccaat	cnangaaacg	nccntgaaag	anccaggcna	anannntccg	780
canatcttat	cccttanttn	ggggnccttt	nccnngggcc	cc		822

<210> 30

<211> 787

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

&lt;222&gt; (1)...(787)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 30

cgccgcctg	ctctggcaca	tgcctvctga	atggcatcaa	aagtgalga	ctgcccatcg	60
ctaggaga	cctctctcc	tactgtratt	atggagccct	gcagactgag	ggctccctt	120
gtctgcagga	tttgatgtct	gaagtcgtgg	agtgtggtt	ggagctcctc	atctacatna	180
gctggagcc	ctggaggggc	tctctcgcca	gctcccccct	tctctcraag	ctctccangg	240
acaccagggg	ctccaggcag	cccattatto	ccagnangac	atggtgttct	tcacgcggga	300
cccatggggc	ctgnaaggcc	agggtctcct	ttgacacccat	ctctcccgtr	ctgcctggca	360
ggcctggga	tcactantt	ctanaacggg	cgccaccncg	gtgggagctc	cagcttttgt	420
tccttttct	gaaggttaat	tgcncgcttg	gcgtatctat	nggtcanaan	tnlttctgt	480
gtgaattgt	ttntccctc	ncnattccnc	ncnacatacn	aaccgggaan	cataaagtgt	540
taaagcctgg	gggtngcctn	ngaatanac	tnaactctaat	taattgcgtt	ggctcatggc	600
ccgtttccn	ttcnggaaa	ctgtctctcc	ctgcnctnnt	gaatcgccca	ccccccnggg	660
aaaagcgggt	tgcctttctg	gggntccctt	ccncttcccc	cctcncctaan	ccctnccgct	720
cggtcgttnc	nggtngcggg	gaaggggnat	nnctcccnnc	naagggggng	agnnngntct	780
cccccaa						787

&lt;210&gt; 31

&lt;211&gt; 799

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(799)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 31

tttttttttt	tttttttggc	gatgctactg	tttaattgca	ggaggtgggg	gtgtgtgtac	60
catgtaccag	ggctattaga	agcaagaagg	agggaggggg	ggcagagrgc	cctgctgagc	120
aacaaaggac	tctgtcagcc	ttctctgtct	gtctcttggc	gcaggccaat	ggggaggccct	180
cccgacaggt	gggggccacc	agtcragggg	tggagguact	acanggggtg	ggagtgggtg	240
gtggctggtn	cnaatggcct	gncacanac	cctacgattc	ltgacacctg	gatttcacca	300
ggggaccttc	tyttctacca	nggnaacttc	ntnnatctcn	aaagaacaca	actgtttctt	360
cngcanttct	ggctgttcat	ggaaagcaca	ggtgtcctat	ltngggtggg	acttggtaga	420
tatggttcgy	gcccacctct	ccntcnaaan	aaqtaattca	cccccccn	ccntctnttg	480
cctgggcctt	taantaccua	cacoggaact	canttantta	ttcatcttng	gntgggcttg	540
ntnatcnccn	cctgaangcg	ccaagttgaa	aggccacgcc	gtncnctc	cccatagnan	600
ntttttnct	cactaatgc	ccccccnggc	aacnatccaa	tcccccccn	tgggggcccc	660
agccacaggg	ccccgctcg	ggnnccngn	cnognantcc	ccaggtctc	ccntcngnc	720
ccnnngcnc	ccgcacgca	gaacanaagg	atngagccnc	cgcannnnnn	nggttncnac	780
ctcgccccc	ccnngng					799

&lt;210&gt; 32

&lt;211&gt; 789

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(789)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 32

tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	60
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	120
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	180
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	240
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	300
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	360
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	420
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	480
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tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	720
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	780
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	789

&lt;210&gt; 33

&lt;211&gt; 793

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{793}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 33

gacaggaacat	gttggatggt	ggagcacctt	tctatacagc	ttacaggaca	gcagatgggg	60
aattcatggc	tgttgagca	atanaacccc	agttctarga	gclgctgato	aaaggacttg	120
gactaaaglc	tpatgaactc	cccaatcaga	tgagcatgga	tgattggcca	gaatgana	180
agaagtittg	agatgtattt	gnaaggaaga	cgaggcaga	gtggatgcaa	atctttgagc	240
gcacagatgc	ctglatgact	cgggttctga	cttttgagga	ggttggtcat	catgatcaca	300
acxangaaag	gggtctggtt	atnaccantg	aggagcagga	ngtgagcccc	cgccctgear	360
ctctgctgtt	aaacaccccc	gcctatccct	ctttcabaag	ggatccacta	cttctagagc	420
gncogccacc	gcggtggagc	tccagctttt	gttcccttta	gtgaggggta	attgagcgct	480
tggcgtaatc	atggtcatan	ctgtttctctg	tgtgaabttg	ttatccgctc	acaattccac	540
acaaacatacg	anccggaage	atnaaatttt	aaagcctggg	ggtngcclaa	tgantgaact	600
nactcanatt	aattgagctt	gcgtcactg	ccgcttttcc	agtcaggaaa	acctgtcctt	660
gcagatgccc	nttaatgaat	cnggccaccc	ccgggggaaa	aggcnqtttg	cttnttgggg	720
ngccttccc	gctttcttgc	ttcttgaant	ccttccccc	ggtctttcgg	cttgaggana	780
acggtatena	cct					793

&lt;210&gt; 34

&lt;211&gt; 756

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{756}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 34

gacaggaacat	gttggatggt	ggagcacctt	tctatacagc	ttacaggaca	gcagatgggg	60
aattcatggc	tgttgagca	atanaacccc	agttctarga	gclgctgato	aaaggacttg	120

ccaaccacag	ggaccaagct	gannaaacag	cagctaattc	tggcccggtga	catactggag	180
atcggggccc	aatggagcat	cctacogcan	gacatcccc	ccttcgagcg	ctacatggcc	240
cagctcaat	gctactact	tgattacaa	gagcagctcc	cagagtcagc	ctatatgrac	300
cagctcttgg	gcctcaaccc	cctcttctg	ctgtcccaga	acggggtggc	tgantccac	360
acgganttgg	aneggctgcr	tgrccaange	calacacacc	aatgtctaca	tunaccacca	420
gtgtcttgg	gcaatattga	tyganngcag	ctaccncaaa	gttttcttgg	ccnagggtta	480
cacccccgc	cgagagctac	accttcttca	ttgaatctct	gtcgcacact	atcagggtatg	540
aaaatcgng	ggtttcttca	gaaaggctnc	aanaaatnc	tttctctga	aggccccgg	600
atnctctagt	ntagaatcg	gccccccatc	gagggtggnc	ctccaacctt	togtttccct	660
ttactgaggg	tttatttgcg	cctttggcgt	tatcatgttc	amucngttn	cctgtattga	720
aattnttaac	cccccaaat	tccagcgcna	cattng			756

<210> 35  
 <211> 834  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (834)  
 <223> n = A,T,C or G

<400> 35						
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tgttcagaca	cncctcttgg	caaaaaacan	caggatntga	gttttgattt	caactccaat	180
aatcttcngg	gctgtctgct	ggttgaactc	gatgacnang	ggcagctggg	tgtgtntgat	240
aaantccanc	angtttctct	tggtagacct	cccttcaag	ttgttcgggc	cttcatcaaa	300
cttctnnaan	angannancc	canctttgtc	gagctggnat	ttgganaaoc	cgtracrgtt	360
ggaaactgat	cccaaatggg	atgtcatcca	tgcctctctg	tgcctgcaaa	aaacttgctt	420
ggcncuaate	cgactcncn	tcttggaaag	aagccnatca	cacccccctc	cctggacttc	480
nncaangact	ctnccgctnc	ccctccnng	cagggttgg	ggcannccgg	gcccctgggc	540
ttcttcagcc	agttcarnat	nttctcagc	ccctctgcca	gctgtnttat	tcttgggggg	600
ggaanccgtc	tctcccttcc	tgaannaart	ttgaccgtng	gaatagccgc	gcnctcncnt	660
acntnctggg	cggggttcaa	antccctccn	ttgncntcn	cctcggggcc	ttctggattt	720
ncnaactttt	ttcttccccc	ccccccnccg	ngtttggntt	tttcatnggg	ccccaaotct	780
gotnttggcc	antccctgg	gggcncttan	ccccccctnt	ggtccctng	ggcc	834

<210> 36  
 <211> 814  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (814)  
 <223> n = A,T,C or G

<400> 36						
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cctagnaaac	attaatgggt	tgtctacta	ataatcata	cnaaccagta	agcctgcccra	120
naacgccaac	tccagccatt	cctaccaaa	gaagaaaggc	tggctctccc	accccctgta	180
ggaaaggcct	gccttctaag	accccaaat	ccggttgaat	ctnaagtctt	gtgttttact	240
aatggssaaa	aaaaataaac	aanaggtttt	gttctctagg	utgcccaccc	cagcctggca	300
ctaaaccanc	ccagcgcctca	cttctgcttg	ganaaatatt	ctttgctctt	ttggacatca	360

ggcttgatgg	tatcaactgcc	acntttccac	ccagctgggc	ccccctccc	catntttgtc	420
antganctyg	agggcctgan	ncttagtctc	caaaagtctc	ngccacacaag	accggccacc	480
aggggagtc	ntttncagtg	gatctgucuu	anantaccn	tatcatcmt	gaataaaag	540
gccccgaac	ganatgcttc	cancancttc	taagaccct	aatccctngaa	ccatggtgcc	600
cttcgggtct	gacccuaaag	gaatgttct	gggtccant	ccctcctttg	tttctacgt	660
tgtnttgga	ccntgctngn	atnacccaan	tganatccc	ngaagraccc	taccctggc	720
atttganttt	cntaaattct	ctgcuctaen	netgaagca	cnatccctn	ggncnnaen	780
ggngaectca	ngaaggtctn	ngaaaaacca	cnen			814

<210> 37  
 <211> 760  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{760}  
 <223> n = A,T,C or G

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gcgagtggt	cgctgaaggy	gttgtagtar	cagcgaggga	tgtctcctt	gcagagtcct	120
gtgtcttgc	ggtccacgca	atgccttttg	tacttgagg	aatggatgg	ctygagtcg	180
tcaanccac	tcgtgtattt	ttccacngca	gctcctccg	agcctccgg	gcagttgggg	240
gtgtcgtcac	actccactaa	actgtcgatn	cancagccca	ttgttgcagc	ggaaactgggt	300
gggtgacag	gtgccagcac	acacttgatn	ggcctttcca	tggagagggc	tgggggaaat	360
ccctnangc	caaaactgct	ctraaaggcc	accttgcaaa	cccgacagg	ctagaaatgc	420
actcttcttc	ctaaaggtag	ttgtctcttg	tggccaagca	ncctccanca	nacaaaaanc	480
ttgcaaaatc	tgtcccttgg	gggtcatnon	taccanggtt	gggaaanaa	accggcngn	540
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caattgaact	gttaacnttg	ggcnggttc	ccctngggctg	gtctgaact	aatcaccttc	660
actggaaaaa	ggtangtgcc	ttccttgaat	tcccaaanct	ccctngnttc	taygtntttt	720
ctcctctacc	ctaaaaatcg	ttttccccc	ccntangggg			760

<210> 38  
 <211> 724  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{724}  
 <223> n = A,T,C or G

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cttccnaaat	tgtccaaacc	cttcnnccaa	atnucattt	ccggggggggg	gttccaaacc	120
caaatataat	ttgganttta	aattaaatnt	tnattngggg	aananaaaaa	atgtnaagaa	180
aatttaaccc	attatnaact	taaatnccn	gaacccttg	gnttccaaaa	atttttaann	240
cttaaatccc	tcogaaattg	ntaanggaaa	accaaattcn	cttaaggctn	tttgaagggt	300
ngatttaaac	ccccctnaat	tttttttacc	cnngnctnaa	ncatttngnt	tccgggtgtt	360
tccntttaan	cntnggtaac	tcccgnatgt	gaanncct	aanccaatta	aacogaaatt	420
tttttgaaat	ggaaactccn	nyggaaattna	ccgggggttt	tccnttttgg	ggggcatncc	480
ccccctttcg	gggtttgggn	ntagggttga	tttttnnang	ccccaaaaaa	ncuccaana	540
aaaaaactcc	caagntttaa	ctngaatntc	cccttccca	ggccttttgg	gaaggngggg	600

tttntggggg cgggggantt ctttccccc tttccccc ccccccnggt aaanggttat 660  
 ngnttttggg ttttggggcc cttttagggc cttccggatn gaattaaat ccccccgggcg 720  
 ggcg 729

<210> 39

<211> 751

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(751)

<223> n = A,T,C or G

<400> 39

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 ttttttttat ttttttttat gttttttttt attttttttt cttttttttt ttttttttat 180  
 gtcggcctta agttttttta atttttttta ttttttttat ctgaanggaa aaggggggtt 240  
 cgcataata cttggggggg nggaaaggtt gtttttttat ttttttttat ttttttttat 300  
 ttttttttat gtttttttat ntttttttat taatttttat ttttttttat ttttttttat 360  
 cttgggggtt ctttttttat accaaccctt ctgaacaaa gtgttcngcc ctttttttat 420  
 ttttttttat ctttttttat ctgttcngaa ngtttttat ntttttttat ctttttttat 480  
 ttttttttat ctttttttat ctttttttat ctttttttat ctttttttat ctttttttat 540  
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<210> 40

<211> 753

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(753)

<223> n = A,T,C or G

<400> 40

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 cgtcttatgc acagctgggc ctttgagaca gtagggcttc gatgtcaggc tctatgttaa 180  
 tctcaaggtt ccaggcaaac tctgttgagc acacccgaga ccagggtgat agcttgggtt 240  
 cgttcataaa cgtgttgagg tctgttgagg gtaggtggag ggttcctcgc aggaaggcna 360  
 ataaagggtg cgttcctggc cgttcctcgc cgttcctcgc cgttcctcgc cgttcctcgc 420  
 cnaaccacc accannccgg atttcttga nggaattccc aaattcttcc gttcttggc 480  
 ttttttttat gtttttttat gtttttttat gtttttttat gtttttttat gtttttttat 540  
 aaanccacc cttctctctt ttttttttat ttttttttat gtttttttat gtttttttat 600  
 gtttttttat ttttttttat ttttttttat ttttttttat ttttttttat ttttttttat 660  
 ttttttttat ttttttttat ttttttttat ttttttttat ttttttttat ttttttttat 720  
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<210> 41

<211> 341  
 <212> DNA  
 <213> Homo sapien

<400> 41  
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 agtgeccccca tcttgattt atatacatat atgttctcag tattttggga gcccttcccac 120  
 ttctttaaac cttgttccct atgaacactg aaatatggaa ttgtgaaga gttaaaaagt 180  
 tatagcttgc ttacgtagtz agtttttgaa gtctacattc aatccagaca cttagttagg 240  
 tgttaactg tgattttta aaatatcat ttgaggaat tctttcagag gtattttcat 300  
 ctttacttt tgattaatg tgttttatat attagggtag t 341

<210> 42  
 <211> 101  
 <212> DNA  
 <213> Homo sapien

<400> 42  
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 gtttcaaaac ttctaataa ataattttca gtggcttcat a 101

<210> 43  
 <211> 305  
 <212> DNA  
 <213> Homo sapien

<400> 43  
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 tccaggttgg tctcacactg taattagagc tattgaggag tctttacagc aatttaagt 120  
 ttagatgctt tgcataagtat agagttctag agttatgttt cagaaagttc aagaaccca 180  
 cctcttgaga ggtcagtaaa gaggaattaa ttttcatat ctacaaaatg accacaggat 240  
 tggatacaga acgagagttc tcttggtata ctcaagagctg agtacctgcc cggggggcgc 300  
 tggaa 305

<210> 44  
 <211> 852  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)... (852)  
 <223> n = A, T, C or G

<400> 44  
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 ctctccatcc tggggcattc ttcccaatt tatataccag tottcgtcca tccacargct 180  
 ccagaatttc tctttttagt taatatctca tagctcggct gagctttcca taggtcatgc 240  
 tgcgttgttt cttcttttta ccccatagct gagccaactgc ctctgatttc aagaaactga 300  
 agacgccttc agatcgggtc tcccaattta ttaatcctgg gttcttgtct gggttcaaga 360  
 ggatgtcggc gatgaattcc cataagtgag tccctctcgg gttgtgtttt ttggtgtggc 420  
 acttgccagg ggggtcttgc tccittttca tatcagggtg ctctgcacaa ggaaggtgac 480  
 tgggtggtgt catgagatc tgagcccggc agaaagtatt gctgtccacc aatctactg 540  
 tgcatacata gttggtgtca tacaatagt cctngtcttt ccaggtgttc atgatggaag 600

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gtcagtttg ttcagtccttg acaatgarat tgtgltgtga ctggaaacagg tcaactactgc      660
actggccgll ccaacttcaga tgcctgcagg tgcctgtagag gaagtgcccc gccgtccctg      720
ccgcccgggt gaactctgc acaactcctgc tgcacagggt ctgcgccgttg atgtcgaact      780
cntggaaagg gatcaattg gcacccagct ggttggtgtc caggaggtga tggagccact      840
cccauacctg gt                                         852

```

&lt;210&gt; 45

&lt;211&gt; 234

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 45

```

acaaacagacc ctitgctcgtt aaagacctca tgcctcatca gttggacgaa tccgtgtccg      60
agttcgacac catccggagc atcagcattg cttegcagtg cctaccgag gggaaatctt      120
gcctcgtttc tggctgggggt ctgctggcga aagggcagaat gctaccgtg ctgcagtgcg      180
tgaacgtgtc ggtggtgtct gaggaggtct gcagtaagct ctatgacccg ctgt      234

```

&lt;210&gt; 46

&lt;211&gt; 590

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(590)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 46

```

actttttatt taaatgttta taaggcagat ctatgagaat gatagaaac atgggtgtga      60
atttgetagc aatatttttg agattacaga gtttttaglaa ttaccaatta caacgttaaa      120
aagagataaa tatattccaa gcaatataca aatatctaat gaagatcaa ggcaggaaaa      180
tgantataac taattgacaa tggaaaatca attttaatgt gaattgcaca ttatcttta      240
aaagctttca aaanaaaaa ttattgcagt ctanttaatt caaacagtyt taaatgggtat      300
caggataaan aactgagggt canaaagaat taattttcac ttcatgtaac ncauccaat      360
ttadaatggc ttaaatgcan ggaanaagca gtggaagtag ggaagtanto aaggtcttct      420
tggctctctaa ctgccttac tctttgggtg tggctttgat cctctggaga cagctgccag      480
ggctcctgtt atatccaca tccagcagc aagatgaagg gatgaagag gacacatgct      540
gccttctctt gaggagactt catctcactg gccaacactc agtcacatgt      590

```

&lt;210&gt; 47

&lt;211&gt; 774

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(774)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 47

```

acaagggggc ataatgaagg agtgggggaa gattttznag aaggaaaaaa aacgaggccc      60
tgaacagaat ttctctgnac aacggggggt caaaataaatt ttcttgggga ggttcagagc      120
gcttcactgc ttgaacctta aatggatgtg ggaacnaatt ttctgtaatg accctgaggg      180
cattacagac gggactctgg gaggaaaggt aaacagaagg gggacaaaagg ctantccaa      240
aacatcaaan aaaggaagggt aggtgcatac ctccuagcct acacagttct ccagggtctt      300

```



```

cttcattcccl: ggaaggacgac agtggaggga ccaactgacca tgcacccagg ctccctgtgtg 360
ctgggtctctg gtcttcagcc cccagctctg gaagccacac ctctgtgat cctgcgtggc 420
ccacactccf. tgaacacaca tccccaggtt atattctctg acatggctga acctctctll 480
cctacttccg agatgctctg ctccctgcag cctgtcaaaa tcccaactac cctcccaacc 540
acggcatggg aagcctttct gacttgcttg attactucag catcttggaa caatccctga 600
ttcccaactc cttagaggca agatagggtg gtttaagagta gggctggacc acttggagcc 660
aggtgtctgg ctcccaattt tggctcattt acgagctatg ggarcttggg caagtatctt 720
tcatttctat gggcctcatt tcttctacac tgcaaaatgg gggataataa tagt 774

```

<210> 48  
 <211> 124  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{124}  
 <223> n = A,T,C or G

```

<400> 48
canaaattga aattttctaa aaaggcatct tctctctata tccataaaal gatataattt 60
ttgcaaatat anagatgtgt cacaatttat aatgttctct aallacagct caacgcaact 120
tggt 124

```

<210> 49  
 <211> 147  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{147}  
 <223> n = A,T,C or G

```

<400> 49
gocgatgcta ctattttatt gaaaggaggt ggggtgttct tattattctc tcaacagctt 60
tgtggctaca ggtgggtgtc gactgcata aaaaattttt tccgggtgat tgcaaaattt 120
ttagggcacc catatcccaa gcaatgt 147

```

<210> 50  
 <211> 107  
 <212> DNA  
 <213> Homo sapien

```

<400> 50
acattaaatt aataaaggga ctgttggggg tctgtcaaaa cacatggctt gatataattgc 60
atgggttagg gttaggagga gttaggcata tgtttcggga gagggtt 107

```

<210> 51  
 <211> 204  
 <212> DNA  
 <213> Homo sapien

```

<400> 51
gtcctaggga gtctagggga cacacgactc tgggttcacg gggcchacac acttgcangy 60

```

cgggagaggaa aggcagagaa glgacacccgt caggggggaaa tyacagaaag gaaatcaag	120
gccttgcaag glcagaaagg ggaactcaggg cltccaccac agccctgccc cacttggcca	180
cctccctttt gggaccagca atgt	204

<210> 52  
 <211> 491  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(491)  
 <223> n = A,T,C or G

<400> 52	
acaaagataa cclttatctt ataacaanaa ttgatagtt tttaaggtta gtattgtgtg	60
gggtattttt cnaagacta asgaqtanc tcaggtaaaa agttagaat gtataaana	120
ccatcagaca ggltttttaa aaacaacata ttacanaatt agacaatcat ccttaaaaaa	180
aaaactctct gtatcaatt ctflcttca aatgactga cttaantatt tttaatatt	240
tcnaaaacac ttctcaaaa atttcaana tggtagcttt canatgtacc ctcagtccca	300
atgttgctca gataaataaa tctcgtgga acttaaccac caccacaagc tttctgggac	360
atgcaacagt gtcttttctc tctttttct tttcttttt ttacaggcac agaaactcat	420
caattttatt tggataaana aggtctcca aattatattg aaaaataaat ccaagttaat	480
atcautcttg t	491

<210> 53  
 <211> 484  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(484)  
 <223> n = A,T,C or G

<400> 53	
acataattta gcagggtca ttacataaag atgctattta ttaanaggtn tatgatctga	60
gtattaarag ttgctgaagt ttggtatttt tatgcagcat tttctttttg ctttgataac	120
actacagaa ccttaaggac actgaanatt agtaagtaaa gtccagaaac attagctgct	180
caatcaaatc tctacataac actatagtaa ttaaaacgtt aaaaaaagt gttgaaatct	240
gcactagtat anaccgtcc tgcaggata anactgctt ggaacagaaa gggaaaaaac	300
agctttgant ttctttgtgc tcatangagg aaaggctgaa ttaccttggt gctctccot	360
aatgattggc aggtcngta antncaaaa catattccaa ctcaacactt cttttcnccg	420
tancctganc ctgtgtattc caggandagg cggatggaat gggucagccc ccggtatctc	480
cant	484

<210> 54  
 <211> 151  
 <212> DNA  
 <213> Homo sapien

<400> 54	
actaaacctc gtgcttgtga actccataca gaaaacggty ccatacctga acacggctgg	60
ccactgggta tactgtgac aaccgcaaca acaaaaacac aatctcttg cactggctag	120
tctatgtcct ctcaagtggc tttttgttg t	151

<210> 55  
 <211> 91  
 <212> DNA  
 <213> Homo sapien

<400> 55  
 acctggcttg tctccgggtg gttcccgagg ccccccacgg tccccagAAC ggacacttcc 60  
 gccctccagt ggatactcga gccaaagtgg t 91

<210> 56  
 <211> 133  
 <212> DNA  
 <213> Homo sapien

<400> 56  
 ggccggatgtg cgttgggttat atacaaatat gtcatttlat gtaagggact tgagtatact 60  
 tggatttttg gtatctgtgt gttgggggga cggtcacgga accaataacc catggatacc 120  
 aagggacac tgc 133

<210> 57  
 <211> 147  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(147)  
 <223> n = A,T,C or G

<400> 57  
 acctctggaga acctgagcgg ctgctccggc tutgggatga ggtgatgcan gcnctggcgc 60  
 gactgggagc tgagcccttc cctttggccc tgcctcagag gattgttggc gaontgcana 120  
 tctcantggg ctggatncat gcagggt 147

<210> 58  
 <211> 198  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(198)  
 <223> n = A,T,C or G

<400> 58  
 acagggatat aggttttag ttattgttat tgcataatan attgaatttt ctgtatactc 60  
 tgattacata ctttctactt ttaaaaaaga tgtaaatctt aatttttatg ccatctatc 120  
 atttacaaat gatttacctt gtaaalgaga agtcattgata gcactgaatt ttaactagtt 180  
 ttgacttcta agtttggg 198

<210> 59  
 <211> 330  
 <212> DNA  
 <213> Homo sapien

&lt;400&gt; 59

acacacaaatg	ggttgtaggg	aaatcattatc	agcaaaaacg	gtgatggcta	ctgaaaagat	60
ccattgaaa	ttatcattaa	tgatttttaa	tgacaaagta	tcacaaaactc	actcaatttt	120
cacctgtgt	agtttgctaa	aatggggagt	aactctagag	caatataagt	atottttgaa	180
tacagtcatt	aaatgacaaa	gccagggcct	acaggtgggt	tcagagcttt	ccagacccag	240
caaaaggaat	ctattttatc	acatgggtct	cgtctgtgac	tcaaaataac	caatgatatt	300
tttctctct	attggacttc	tttgaagagt				330

&lt;210&gt; 60

&lt;211&gt; 175

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 60

accgtgggtg	ccttctacat	tctgacggc	tccttcacca	acatctgggt	ctacttggc	60
gtcglgggt	ccttctctt	catctcctc	cagctgggtg	tgctcatcga	cttgcgac	120
tcttggaaac	agcgtggt	gggcaaggc	gaggaagtgc	attccgtgc	ctggc	175

&lt;210&gt; 61

&lt;211&gt; 154

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 61

acccacttt	tcttctgtg	agcgtctgg	acttctcct	gtacatgat	gagggtagt	60
ggttggtgt	cttcaacgt	atctccctt	ttccggatct	gtgagcggg	acagcagtg	120
tggactgcac	agccccggg	ctccacattg	ctgt			154

&lt;210&gt; 62

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 62

cgtctgagc	ctatagttag	tcttattag				30
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&lt;210&gt; 63

&lt;211&gt; 89

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 63

acaagtctt	tcagacctt	ttgctctta	aaactgacca	tcttttatat	ttaatgtctc	60
ctgtatgaat	aaaaatggt	atgtcaagt				89

&lt;210&gt; 64

&lt;211&gt; 97

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 64

acgggagtaa	ctagtgagg	acgtggaac	tgaatccacc	aataaataaa	ggttctgcag	60
aatcagtgca	tccaggattg	gtcttggat	ctgggt			97

<210> 65  
 <211> 377  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (377)  
 <223> n = A,T,C or G

<400> 65  
 acaacaanaa ntcccttctt taggccaactg atggaaacct ggaacccctt ttgatggca 60  
 gcctggcgct ctaggcttg acacagcggc tggggtttgg gctntccraa accgcacacc 120  
 ccaacccctg tctacccaca ntcttggtta tgggtgtgt ctgcactga acatcagggt 180  
 tgggcatata natgaaatcc caanggggac agaggtcagt agaggaagct caatgagaaa 240  
 ggtgtgttt gctcagccag aacacagctg cctggcattc ggcgtgaa cttgaacccg 300  
 tgggggtgaa ctaccccccag gaggaaatcat gctggggcga tgganggtg ccaacaggag 360  
 gggcgggagg agcctgt 377

<210> 66  
 <211> 305  
 <212> DNA  
 <213> Homo sapien

<400> 66  
 acgcctttcc ctacgaattc agggaaagaga ctgtcgcttg ccttccctcg ttgttgctg 60  
 agacccgtg tgcctcttcc caacatatac acccttgcct catctttgaa ctcaaacag 120  
 aggaactaac tgcacctgg tctcttcccc agtccacagt tcaacotcca tccctcact 180  
 tcttccctc taagggatct caacactgac cagacacagg gccctgaatt tatgtggttt 240  
 ctatatattt tttaataaga tgcacttat gtcatttttt aatcaagtct gaagaattac 300  
 tgttt 305

<210> 67  
 <211> 385  
 <212> DNA  
 <213> Homo sapien

<400> 67  
 actacacaca ctccacttgc ccttgtgaga caatttgtcc cagcacttta ggaatgctga 60  
 ggtcggacca gccacatctc atgtgcaaga ttgccagca gacatcaggt ctgagagttc 120  
 cctttttaaa aaaggggact tgettaaaaa agaagtctag ccargatgt gttaggcagc 180  
 tgtgtgtgtc tgyagattca cttttgagag agttctctc tgaagacctga tctttagagg 240  
 ctgggcagtc ttgcacatga gatggggtg gtctgatctc agcactcctt agtctgcttg 300  
 cctctccag ggcaccagc tggccacac tgettacagg gcactctcag atgccatcc 360  
 catagtttct gtgctagtgg accgt 385

<210> 68  
 <211> 73  
 <212> DNA  
 <213> Homo sapien

<400> 68  
 acttaaccag atacatttt accccagatg gggatattct ttgtaaaaa tgaanataaa 60  
 gtttttttaa tgg 73

<210> 69  
 <211> 536  
 <212> DNA  
 <213> Homo sapien  
 <220>  
 <221> misc\_feature  
 <222> (1)... (536)  
 <223> n = A,T,C or G

<400> 69  
 actagtcacag tgtgggtggaa ttccattgtg ttggggggtc tcacctctct ctctgcagc 60  
 tcacgctttg tgcctctgct ctgaggagag catgcccag catctgagta ccttgcctgt 120  
 cctgctggcc accctagctg tggcctggc ctggagcccc aaggaggagg ataggataat 180  
 cccgggtggc atctataac cagacntcaa tcatgagtg gtacagcgtg cctttcactt 240  
 cyccatcagc gagtataana aggcacccaa agatgactac tacagacgtc cyctgcgggt 300  
 actaagagcc aggcacacga ccgttggggg ggtgaattac ttcttcgacg tagaggtggg 360  
 ccgaaccata tgtaccaagt cccagcccaa ctgggacacg tgtgcttcc atgaacagcc 420  
 agaactgcag aagaaacagt tgtgctcttt cgaatctac gaagttccct ggggagaaca 480  
 gaangtccct gggtgaaatc caggtgtcaa gaaatcttan ggatctgttg ccaggc 536

<210> 70  
 <211> 477  
 <212> DNA  
 <213> Homo sapien

<400> 70  
 atgaccccta acagggggcc tctcagccct cctaattgac tcgggctag cctgtgtgatt 60  
 teacttccac tccctaacgc tctctactt aggcctacta accaacacac taacctata 120  
 ccgatgatgg ccgatgttaa caggagaaag cactatccaa ggcacccaca cccacccgt 180  
 ccnaaaagcc ctctgataag ggataatcct atttattacn tcagaagttt tttctctgc 240  
 agggattttt ctgagccttt taacactcva gcttagcccn taccccccaa ctaggagggc 300  
 actggccccc aacaggcatc cccccgctaa atnccctaga agtcccactc ctanacacat 360  
 ccgtattact cyccatcagg gtatcaatca cctgagctca ccatagtcta atagaaaca 420  
 accgaaccca aattattcaa agcactgctt attacaattt tactgggtct ctatttt 477

<210> 71  
 <211> 533  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)... (533)  
 <223> n = A,T,C or G

<400> 71  
 agagcttatg gtacagtgtg atctcagrtt tgcacacaca tttttctacat agatagtact 60  
 aggtattaat agatatgtaa agaaagaaat cacaccatta ataatggtaa gattgggtta 120  
 tgtgatttta gtggtatttt tggcaccrtt atatatgttt tccaaacttt cagcagtgat 180  
 attatttcca taacttaaaa agtyagtttg aaaaagaaaa tctccagcaa gcatctcatt 240  
 taaataaagg tttgtcatct ttaaaaatcc agcantatgt gactttttta aaaagctgtc 300  
 aaataggtgt gacctacta ataattatta gaalacatt taaaaacatc ggtacctca 360  
 agtcagtttg ccttgaaaaa tatcaaatat aactcttaga gaaatgtaca taaaggaatg 420  
 cttcgttaatt ttggagtang aggttccttc ctcaattttg tatctttaa aagtacatgg 480  
 taaaaaaas eattccacac agtatataag gctgtaaaat gaaagattct gcc 533

<210> 72  
 <211> 511  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(511)  
 <223> n = A,T,C or G

<400> 72  
 tatctacggaa aaacacacca cataattcaa ctanccaaaga anactgcttc agggcgtgta 60  
 aatgaaagg cttccaggca gttatctgat taaagaacc taaagagggg acaaggctaa 120  
 aagccgcagg atgtctacac tatancaggc gctatttggg ttggctggag gactgtgga 180  
 aacratggan agattggtgc tgganacgc cgtggctatt cctcattgtt attacanagt 240  
 gaggttctct gtgtgccac tggtttgaan accgttctnc aataatgata gnatagtaca 300  
 cacatgagua ctgaaatggc ccaaacuccg aaagaaagcc caatagatc ctragaannac 360  
 gcttctaggg acaataaccg atgaagaaa galggctctc ttgtgcccc gtctgttatg 420  
 atttctctcc attgragcna naaacccgtt cttctaagca aacnucggtg atgatggcna 480  
 aaatacaacc cctcttgaag naccnggagg a 511

<210> 73  
 <211> 499  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(499)  
 <223> n = A,T,C or G

<400> 73  
 cagtgcacgc actggtgcca gtaccagtae caataacagt gccagtacca gtgcacagcac 60  
 cagtggatgg ttcagtgttg gtgccagccg gccggccact ctacatcttg ggcctcttgc 120  
 tggccttggg ggagctggat ccagcaccag tggcagctct ggtgactgtg gtttctctca 180  
 caagtgaat tttagatatt gttaatcttg ccagtcttct tcttcaagcc aggggtgcac 240  
 ctacagaaac tactcaacac agcaactetg gcagccacta tcaatcaatt gaagctgaca 300  
 ctctgcatta atctatcttg ccatttctga aaaaaaanaa aaaaaagggg cggccgcttg 360  
 antctagagg gccctgttaa acccgtgat cagcctcgac tgtgcttct anttgcacgc 420  
 catctgttgt ttgccctcc ccgntgcct tcttgaccc tggaaagtgc cactccact 480  
 gtcccttctc aataaact 499

<210> 74  
 <211> 537  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(537)  
 <223> n = A,T,C or G

<400> 74  
 tttcataggg gaacacactg agagatact ttaagaattt ggaatcagcc gcaagagat 60





```

ccagctgccc cggcggggga tgcgaggctc ggagcaccct tgcggggctg tgattgctgc      120
caggcactgt tcatctcaga tttctgtcc ctttgcctcc ggcagcgctc tctgtgaaa      180
gttcatatct ggagcctgat gtcttaacga ataaaggctc catgctccac ccgaacaaa      240
aaaaaaaaa                                     248

```

<210> 78  
 <211> 201  
 <212> DNA  
 <213> Homo sapien

```

<400> 78
actagtcacg tgcggtgga ttccattgtg ttcgggcaca cacaatggct acctttaaca      60
tcacccagac ccgcctctgc cngtgcacca cgtgctgctc aacgacagta tcatgcttac      120
tctgctactc ggaaacatct tttatgtaac taatgtatgc tttcttgctt ataatgcct      180
gatttaaaa aaaaaaaaaa a                                     201

```

<210> 79  
 <211> 552  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (552)  
 <223> n = A,T,C or G

```

<400> 79
tctttttgtt aggtttttga gacaaccctc gacctaaact gtgtcacaga cttctgaatg      60
tttaggcagt gctagtaatt tctctgtaat gattctgtta ttactttctt attctttatt      120
cctctttctt ctgaagatta atgaagttgc aaattgaggt ggataaatat aaaaaggtag      180
tgtgatagta taagtatctc agtgcagatg aaagtgtgtt atatatactc attcaaaatt      240
atgcaagtta gtaattactc aggtttaact aaattacttt aatatgctgt tgaacctart      300
ctgttccttg gctagaaaaa attataaaca ggaactttgtt agtttgggaa gccaaattga      360
taatattcta tgttttaaaa gttgggctat acataaanta tnaagaataa tgggaattta      420
tcccaggaa tatggggttc atttatquat antaccggg anagaagttt tgantnaaac      480
cngttttggg taatacgtta atatgtcttn aatnaacaa gctgactta ttccaaaaa      540
aaaaaaaaa aa                                     552

```

<210> 80  
 <211> 476  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (476)  
 <223> n = A,T,C or G

```

<400> 80
acagggattt gagatgctaa ggcccagag atcgtttgat ccaacctct tttttcaga      60
ggggaaaaatg gggcctagaa gttacagagc atctagctgg tgggttggca cccctggcct      120
cacacagant ccgaggtagc tgggactaca ggcacacagt cactgaagca ggccctgttt      180
gcaattcaag ttgccacctc caacttaaac attcttcata tctgatgtcc ttagtcacta      240
aggttaaac ttcacaccca gaaaggcaa cttaagataa atcttagagc accttcatac      300
tcttttaagt ccttttcacg cctcacttct agtctcctt gggggttgat aggaantctc      360

```

tcttgggtt1 ctcataaana tctctatcna tctcatgttt aatttgggtac gcntaaaaat 420  
 gctgaanaaa ttaaatgtt clggttttnc tttaaaaaa aaanaaaaaa aaaaaa 476

<210> 81  
 <211> 232  
 <212> DNA  
 <213> Homo sapien  
 <220>  
 <221> misc\_feature  
 <222> (1)... (232)  
 <223> n = A,T,C or G

<400> 81  
 tttttttttg tatgccttcn ctgtggngtt attgttgcg craccctgga ggagcccagt 60  
 ttctttctgt ttttttttt ctgggggato ttcttggtc tgcacctera ttcccagcct 120  
 ctcatccca tcttgcaatt ttgttaggtt tggagggcgt ttcttggtag cccctcagag 180  
 actcagtcag cgggaataag tccatagggg ggggggtgtg gcaagccggc ct 232

<210> 82  
 <211> 383  
 <212> DNA  
 <213> Homo sapien  
 <220>  
 <221> misc\_feature  
 <222> (1)... (383)  
 <223> n = A,T,C or G

<400> 82  
 aggcggggagc agagcctaaa gccaaagccc aggaagagtg gcagtgcag caetgggtgc 60  
 agtaccagta ccaataacat gccagtgcc gtcaccagcac cagtgggtgc ttcaagtgcg 120  
 gtgccagcct gaccgcaact ctacacattg ggtctctgc tggccttggg ggagctggg 180  
 ccagcaaccag tggcagctct ggtgctgtg gtttctccta caagtggat tttagatatt 240  
 gtttaactctg ccagtcttct tttcaagcc aggytgcac ctcaagaacc tactcaaac 300  
 agcactctng gcagcacta tcaatcaatt gaagttgaca ctctgcatta aatctattg 360  
 ccatttcnaa aaaaaaaa aaa 383

<210> 83  
 <211> 494  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)... (494)  
 <223> n = A,T,C or G

<400> 83  
 accgaatttg gaccgctggc ttataagoga tcatgtctc cagtattacc tcaaggagca 60  
 gggagatcga gtctatacgc tgaagaaatt tgaccogtg ggacaacaga cctgtcagc 120  
 ccattctgct cggttctccc cagatgaca atactctga caccgaatca ccatcaagaa 180  
 acgttcaag gctgtcatga ccagcaacc ggcacctgc ctctgagggg ccttaactg 240  
 atgtctttt tgcacctgt taccctctg agactcagta accaactct tgggactgt 300  
 agccctgatg cctttttgac agccatactc ttgggentec agtctctcgt ggcgattgat 360

```

talgcttctg  ttaggcacac atggtggcat caccacatnaa gggaacacat ttgattttt 420
tttncatat  tttaatatc nccagaaata nttragaata aatgaattga aaacactta 480
aaaaaagaaa  aaaa 494

```

```

<210> 84
<211> 380
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(380)
<223> n = A,T,C or G

```

```

<400> 84
gctggtagcc tatggcgtgg ccacggaggg gctcctgagg cacgggacag tgacttccca 60
agtatacctgc gccgcgtctt ctaccgtccc tccctgcaga tcttggggca gattccccag 120
gaggacatgg acgtggccct catggagcac agcaactgct cgtcggagcc cggcttctgg 180
gcacacccctc ctggggccca ggccggccac tgcgtctccc agtatgccaa ctggctgggtg 240
gtgctgctcc tctcatctt cctgctcgtg gccaacatcc tgcctgctcc ttgctcctg 300
ccatgttcag ttaacacctc gccaaagtac agggcaacag cnatctctac tgggaaggcc 360
agcgttcccy cctcatccgg 380

```

```

<210> 85
<211> 481
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(481)
<223> n = A,T,C or G

```

```

<400> 85
gagttagctc ctccacaacc ttgatgaggt cgtctgcagt ggctctctgc ttcataccgc 60
tncatcgtc atactgtagg ttbgccacca cctcctgcat cttggggcgg ctaatatcca 120
ggaaactctc aatcaagtea ccgtcnatna aacctgtggc tggttctgtc tccgctcgg 180
tgtgaaggga tctccagaag gaggctctga tcttccccac acctttgatg actttattga 240
gtcgaattctg catgtccagc aggaggttgt accagctctc cgacagtga gtcaccagcc 300
ctatcatgc nttgaacgtg ccgaayaaca ccgagccttg tgtggggggt gnegtctcac 360
ccagattctg cattaccaga naccgtggc aaaganatt gacaaactgc ccaggngaa 420
aaagaacacc tcttggaggt gctngccgt cctcgtcctt tgggtggangc gcntaccttt 480
t
481

```

```

<210> 86
<211> 472
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(472)
<223> n = A,T,C or G

```

```

<400> 86

```

aacatcttcc	tgtataatgc	tgtgtaatat	cgatccgagc	ttgtctgctg	agaattcatt	60
acttggaaa	gaaacttnaa	gcctgggacg	tggtattaaa	attcacaala	tgcaacactl	120
taaacagtgt	gtcaalctgc	tcccttactt	tgtatccacc	agtctgggaa	taaggggtatg	180
cctattccg	acctgttaaa	agggcgclaa	gcatttttga	ttcaacatcl	ttttttttga	240
cacaagtcg	aaaaaagcaa	aagtaaacag	ttnttaattl	gttagccaat	tcactttctt	300
cctgggacag	agccatttga	tttaaaaaag	aaattgcata	atattgagcl	ttgggagctg	360
atatntgagc	gggaagatcg	cctttctact	tcaccagaca	caactccttt	catattggga	420
tgttaacaa	agtcattgtc	cttaccagatg	ggtatgcttt	gtggcaattc	tg	472

<210> 87  
 <211> 413  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(413)  
 <223> n = A,T,C or G

agaaaccagt	atctctnaaa	acaaacctctc	atacccttgc	gacctaatct	tgtgtgagtg	60
tgtgtgtgag	agcatattat	atagacaggc	acatcttttt	tactttttga	aaagcttatg	120
cctcttttgg	atctatatct	gtgaaagtct	taattgatctg	ccataatgct	ttggggacct	180
ttgtcttctg	gtlaaatggg	actagagaaa	acacctctct	tatgagtcaa	tctagttingt	240
tttattcgac	atgaaggaaa	tttccagatn	acaaactttn	caaaactctcc	cctgactagg	300
ggggacaaa	aaaagcnaa	ctgaacatna	gaaacaattn	cctggttgaga	aatttncataa	360
acggaaattg	ggtngtatat	tgaaanang	catcattnaa	acgttttttt	ttt	413

<210> 88  
 <211> 448  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(448)  
 <223> n = A,T,C or G

cgcagcgggt	cctctctatc	tagctccagc	ctctcgctcg	ccccactccc	cgcgtccgc	60
gtcctagccn	accatggccg	ggcccttggc	cgcctccgctg	ctctgctgg	ccatcctggc	120
cgtggccctg	gccgtggccc	ccgcggcccg	ctccagtcct	ggcaagccgc	cgcgcctgg	180
gggaggccca	tgyaccrcgc	gtggaagaag	aaggtgtgag	gcgtgcactg	gactttgccc	240
tggcnanta	caacaaaccc	gcacnaactt	ttaccnagcn	cgcgtctgcg	gttctgccc	300
ccccancaa	ttgttactng	gggtaantaa	ttcttgyaag	ttgaacctgg	gucnaacnng	360
tttaccagaa	ccnagccaat	tngaacaatt	ccccctccat	aacagccctt	tttataaagg	420
gaancantcc	tgntcttttc	caaatttt				448

<210> 89  
 <211> 463  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature

&lt;222&gt; (1)...(463)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 89

gaattttgtg cactggccac tgtgatggaa ccattggggc aggatgcttt gaggtttatca	60
gtagtgattc tgcuaaagtt ggtgttgraa cabgagtatg taaaatgtca aaaaatttagc	120
agaggcttag gtctgcatac caggagacag tttgtcngtg tattttttag ccttgaagtt	180
ctcagtpaca agtttnttct gatgcgaagt tctnattcca gtgttttagt cctttgcate	240
tttnatgtn agacttgcct ctatnaaatt gcttttgnt tctgcaggta ctatctgttg	300
tttaacaaa tagaannact tctctgcttn gaanatttga atatcttaca tctnaaaatn	360
aattctctcc ccatannaaa acccangccc ttggganaat ttgaaaaang gntccttern	420
aattcnana anttcagtn tcatacaaca naaunggan ccc	463

&lt;210&gt; 90

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(400)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 90

agggattgaa ggtctntnt actgtcggac tgttcacca ccaactctac aagttgctgt	60
cttccartca ctgtctgtac gcatnttaac ccagactgta tcttcataaa tagaacaatt	120
tcttcaccag ccacatcttc taggaccttt ttggattcag ttagtataag ctcttccact	180
tcctttgtta agacttcate tggtaaaagtc ttaagtttfg tggaaaggaa tttcaattgct	240
cgttctctaa caatgtcttc tccttgaagt atttggctga acaaccacac tnaagtcctt	300
ttgtgcaccc attttaata cacttaatag ggcattggtn cactaggtta aattctgcaa	360
gagtcacttg tctgcacaaag ttgcgttagt atatctgcca	400

&lt;210&gt; 91

&lt;211&gt; 480

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(480)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 91

gagctoggat ccaataatct ttgtctgagg gcagcanaca tatncagtgc catggnaact	60
ggtctacccc acatgggagc agcatgcctt agtatataaa ggtcattccc tgagtcagac	120
atgctctttt gactaccgtg tgcagtgctt ggtgattctc acacaacctcc nncgctctt	180
tgtggaaaaa ctggcacttg nctggactca gcaagaratc acttacaaat tcaccacaga	240
garacttga aaggtgtaca aagcgactct tgcattgctt tttgtccctc cggcaccagt	300
tytcaatact aacccgctgg tttagctcca tccatttgtt gatctgttag tctggataga	360
tctactgaca gtaetgaaga acttcttctt ttgtttcaaa agcgaactctt ggtgctgtt	420
ngatcagggt cccatttccc agtcogaatg ttcacatggc atatnttact tcccacaaa	480

&lt;210&gt; 92

&lt;211&gt; 477

&lt;212&gt; DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(477)

<223> n = A,T,C or G

<400> 92

ataacagccca	natcccacca	ogaagatgag	cttgttgact	gagaacctga	tcgggtcact	60
ggccccgtg	tggccccagc	gactctccac	ctgctggagc	cggttgatgc	tcgactcctt	120
ccacgcagg	cagcagcggg	gcgggtrcat	gaactccact	cgtgggttgg	ggttgacggg	180
taantgcagg	aagaggctga	ccacctcgcg	gtccaccagg	atgcccgaat	gtgcgggacc	240
tgcagcgaaa	ctcctcgatg	gtcatgagcg	ggaagcgcat	gangccuagg	gccttcacca	300
gaaccttcgg	ctgtttctct	ggcgccacct	gcagctgctg	ccgctnacac	tgggctcggg	360
accagcgggc	aaacggcggt	gacacgcccc	acctccaggc	tgcacantgt	gtngcgtcc	420
aggaacggcg	ccagcgtgtc	caggtccatg	tgggtgaanc	ctccgcgggt	aatggcg	477

<210> 93

<211> 377

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(377)

<223> n = A,T,C or G

<400> 93

gaacggctgg	accttgccct	gcattgcgct	gttggcagga	ctaccttggc	aagcagctcc	60
agtccgaqda	gccccagacc	gctgcggccc	gaagctaaac	ctgcctctgg	ccttcacctc	120
cgctccaatg	cagacccant	agtgggagca	ctgtgtttag	agctaugagt	gaacactgtc	180
tgattttact	tgggaatttc	ctctgttata	tagcttttcc	caatgctaat	ttccaaacaa	240
caacacacaa	ataacatggt	tgcctgttnc	gttgtataaa	agtcngtgat	tctgtatnta	300
aagaaaatal	tactgttaca	tatactgott	gcaanttctg	tatttattgg	tcctctggaa	360
ataaatatat	tatttaa					377

<210> 94

<211> 495

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(495)

<223> n = A,T,C or G

<400> 94

ccctttgagg	ggttagggtc	cagttcccag	tgggaagaaac	aggucaggag	aantgcgtgc	60
cgagctgang	cagatttccc	acagtgaccn	cagagccctg	ggctatagtc	tctgacctct	120
ccaaggaaag	accaccttct	ggggacatgg	gctggaggcc	aggacctaga	ggvaccaagg	180
gaaygcccca	ttccgggggt	gttccccgag	gaggaaggga	aggggtctctg	tgtgcovccc	240
acgaggcaaa	ggccttgant	ctcgggatac	nacacccctt	cacgtgtatc	ccacacacaa	300
tguagctca	ccagggtccc	ctctcagtc	cttccctaca	ccctgaacgg	ncactggccc	360
acacccccc	agancancca	ccgcacatgg	ggaatgttct	caaggatcga	ungggcaacg	420
tggactctng	tccnnnaagg	gggcagaatc	tccaatagan	gganagaacc	cttgcctana	480

AAAAAAAAAAAA

495

<210> 95  
<211> 472  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> {1}...{472}  
<223> n = A,T,C or G

<400> 95  
ggttacttgg tticattgac accacttagt ggatgtcatt tagaaccatt ttgtctgctc 60  
cctctggaag ccttgcgcag agcggacttt gtaattgttg gagaataact gctgaatttt 120  
tagctgtttt gacttgatc gcaccactgc accacaactc aatatgaaaa ctatttnact 180  
tatttattat cttgtgaaaa gtatacaatg aaatttttgt tcatactgta ttatcaaagt 240  
atgatgaaaa gcaatagata tatattcttt tattatgttn aattatgatt gccattatta 300  
atcggcgaag tctggagctg atgtttcttt cacagtaata tatgcctttt gtaacttcac 360  
ttggttattt tatctgtaat gaattacaaa attottaatt taagaagatg gtangtcata 420  
ttanttcacn taatttcttt cttgttttac gtaattttt aagaagatgc at 472

<210> 96  
<211> 476  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> {1}...{476}  
<223> n = A,T,C or G

<400> 96  
ctgaagcatt tcttcaaac tntctacttt tgtcattgat acctgtagta agttgacaat 60  
gtggcgaaat ttcaaaatta tatgtaactt ctactagttt taatttctcc cccaagtctt 120  
tttaactca tgatttttac acacacaaac cagaacttat tatatagcct ctgaagtctt 180  
attcttcaca gtagatgatg aaagagtcct ccagtyctct gngcnaatg ttctagnctt 240  
agctggatac ataungtggg agttctctaa actcatacct cagtgaggact naaccanaat 300  
tgtgttagtc tcaattctta ccaactgag ggagctctcc aatcactat attcttatct 360  
gcaggtactc ctocagaaaa acngacaggg caggtctgca tgaanaagtn acatctgcgt 420  
tcacaagtct atcttctca nangtctgtn aaggaaacat tcaatcttct agcttt 476

<210> 97  
<211> 479  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> {1}...{479}  
<223> n = A,T,C or G

<400> 97  
actctttcta atgctgatct gatcttgagt ataagaatgc atatgtcact agautggata 60  
aaataatgct gcaaaactaa tttcttatg caaantggaa cgtcaatgaa acacagctta 120

caatcgcaaa	tcaaaactca	caagtgcctc	tctgttgtag	atttagtgta	ataagactta	180
gcltgggclc	cltcggatct	gattgtttct	canatcttgg	gcaatntlcc	ttagtcaaat	240
caggctacta	gaattctgtt	attggatctn	tgagagcatg	aaatttttaa	naatcacatt	300
gtgattatna	aattaatcuc	aaatttcact	tatacutgct	atragcagcl	agaaaaacat	360
ntntttttta	natcaaatga	tcttctgttt	ggaantgttn	aaatgaaatc	tgaatgtggg	420
ttenatctta	ttttttccen	gacnaactant	tactttttta	gggactcttc	tyanccatc	479

&lt;210&gt; 98

&lt;211&gt; 461

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 98

agtgaattgt	cctccaaacaa	aaccctttga	tcaagtttgt	ggcactgaca	atcagaccta	60
tgttagttcc	tgtcatctat	tggctactaa	atgcagactg	gaggggacca	aaaaggggca	120
tcaactccag	ctggatttatt	ttggagcctg	caaatctatt	ctactttgla	vggactttga	180
agtgtattcag	tttctctctac	ggatgagaga	ctggctcaag	aatatcttca	tgcagcttta	240
tgaagccact	ctgaacacgu	tggttatcta	gatgagaaac	gagaaatcaa	gtcagaaaak	300
ttacctggag	aaaagaggct	ttggctgggg	accatcccat	tgaaaccttct	cttgaaggact	360
ttaagaaaaa	ctaccacatg	ttgtgtatcc	tggtagccgg	cgtttatgaa	ctgaccaccc	420
tttgaataaa	tcttgargct	cctgaacttg	ctctcttgag	a		461

&lt;210&gt; 99

&lt;211&gt; 171

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 99

gtgggcgcgc	gcagggtgtct	ctctgtaccg	caggggcacc	tcccttcccc	aggcgtccct	60
cggcgctctt	gggggcccga	ggaggagcgg	ctggcggttg	gggggagtgt	gaccaccccl	120
cgggtagaaa	agccttctct	agcgatctga	gaggcgtagc	ttgggggtac	c	171

&lt;210&gt; 100

&lt;211&gt; 269

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 100

cggcgcgaag	tgcaactcca	gctggggccg	tgcggacgaa	gattctgcca	gcagttggto	60
cgaactgcga	gacggcggtg	gcgacagtcg	caaggtgcagc	gcgggcgcct	ggggctcttc	120
aaggttgagc	tgacgcccga	gaggtcgtgt	cacgtcccac	gacrttgacg	ccgtcgggga	180
cagccgggaa	agagcccggt	gaagcgggag	gcctcgggga	gccccctcgg	aagggcggcc	240
cgagagatac	gcaggtgcag	gtggccgc				269

&lt;210&gt; 101

&lt;211&gt; 405

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 101

tttttttttt	ttttggaatc	tactgagagc	acagcaggtc	ayuaacaagt	ttattttgaa	60
gctagcaagg	tacacgggta	gggcattggt	acatgtttcag	gtcaacttcc	tttgtcgtgg	120
ttgattgggt	tgtctttatg	ggggcggggt	ggggtagggg	aaacgaagca	aataacatgy	180
agtgggtgca	cctccctgtg	agaacctggg	lacaagcttt	ggggcagttc	acctgggtctg	240
tgaacctcat	ttctttgaca	tcaatgtttt	tagaagtccg	gntatctttt	agagagtcac	300



ctgttcttga gggagattag ggtttcttgc caaatccaac aaaaatccact gaasaagtgt 360  
 gctgatcagt acgaataccg aggcataatc tcatatcggt ggcca 405

<210> 102

<211> 470

<212> DNA

<213> Homo sapien

<400> 102

tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt 60  
 ggcacttaat ccatattttat ttcaaaatgt ctarcaaattt aatccccatta tacgggtattt 120  
 tcaaaatcta aattattcaa attagccaaa tccctaccac aatatcccc aaaaacaaa 180  
 atatacttct ttcagcaaac ttgttacata aattaaaaaa atatatcagg ctgggtgtttt 240  
 caaagtacaa ttatcttaac actgcaaac ltttaaggaa ctaaaataaa aaaaaaactt 300  
 ccgcaagggt taaagggaac aacuaattct ttaracac cttatataaa atcatatctc 360  
 aatctttagg ggaatatata cttcacacgg gatcttaact ttacttact ttgtttattt 420  
 ttttaacca ttgtttgggc ccaacacaaat ggaatcccc ctggactagt 470

<210> 103

<211> 581

<212> DNA

<213> Homo sapien

<400> 103

tttttttttt ttttttttga cccccctctt ataaaaaaca agttaccatt ttattttact 60  
 tacacatatt tattttataa ttggtatttag atattcaaaa ggcagctttt aaatccaac 120  
 taaatggaaa ctgccttaga tacataattc ttgggaatta gcttaaaatc tgcctaaagt 180  
 gaaaatcttc tctagctctt ttgactgtta atttttgact ctgttaaaac atccaaattc 240  
 atttttcttg tctttaaat tatctaactt ttccattttt tccctattcc aagtcatttt 300  
 gcttctctag cctcatttcc tagctcttat ctactattag taagtggttt ctttctaaa 360  
 agggaaaaca ggaggaqaaa cggcacacaa aacaaacatt ttatattcat atttctact 420  
 acgttaataa aatagcattt tgtgaagcca gtcanaaga aggcctagat ctttttatgt 480  
 ccatttttagt cartaaacya ttcanaagt cagaatgca aaaggtttgt gaacatttat 540  
 tcaaaagcta atataagata ttccanatac tcatctttct g 581

<210> 104

<211> 578

<212> DNA

<213> Homo sapien

<400> 104

tttttttttt tttttttttt tttttctctt cttttttttt gaaatgagga tcgagttttt 60  
 cactctctag atagggcatt aagaaaactc atctttccag ctttaaaata acaatccaat 120  
 ctcttatgot atatcatatt ttgaagttaa ctaatgagtc actggcttat ctctcctga 180  
 aggaatctg ttcattcttc tcatcatat agtlatatca agtactacct tgcattattga 240  
 gaggtttttc tctctctatt acacatatat tccatgtga atttgtatca aacctttatt 300  
 ttcatgcaaa ctagaaaata atgtttcttt tgcataagag aagagaacaa tatagcatte 360  
 caaaactgct caaattgttt gtttaagttat ccattataat tagttggcag ggcctaatat 420  
 aaatcacatt tarcacagca ataatcaaac tgaagtacca gttcaaatat caaaataatt 480  
 aaagggaact ttttagctg ggtataattt gctaatctac tttaacagca tttattagaa 540  
 tgaattcaaa tgttattatt cctagcccaa cacaatgg 578

<210> 105

<211> 538

<212> DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 105

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&lt;210&gt; 106

&lt;211&gt; 473

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 106

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&lt;210&gt; 107

&lt;211&gt; 1621

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 107

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a 1621

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&lt;210&gt; 108

&lt;211&gt; 382

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 108

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20 25 30
Arg Val Asp Arg Pro Gly Ser Arg Tyr Asp Val Ser Arg Leu Gly Arg
35 40 45
Gly Lys Arg Ser Leu Val Leu Asp Leu Lys Gln Pro Arg Gly Ala Ala
50 55 60
Val Leu Arg Arg Leu Cys Lys Arg Ser Asp Val Leu Leu Glu Pro Phe
65 70 75 80
Arg Arg Gly Val Met Glu Lys Leu Gln Leu Gly Pro Glu Ile Leu Gln
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Arg Glu Asn Pro Arg Leu Ile Tyr Ala Arg Leu Ser Gly Phe Gly Gln
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Ser Gly Ser Phe Cys Arg Leu Ala Gly His Asp Ile Asn Tyr Leu Ala
115 120 125
Leu Ser Gly Val Leu Ser Lys Ile Gly Arg Ser Gly Glu Asn Pro Tyr
130 135 140
Ala Pro Leu Asn Leu Leu Ala Asp Phe Ala Gly Gly Gly Leu Met Cys
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Ala Leu Gly Ile Ile Met Ala Leu Phe Asp Arg Thr Arg Thr Asp Lys
165 170 175
Gly Gln Val Ile Asp Ala Asn Met Val Glu Gly Thr Ala Tyr Leu Ser
180 185 190
Ser Phe Leu Trp Lys Thr Gln Lys Ser Ser Leu Trp Glu Ala Pro Arg
195 200 205
Gly Gln Asn Met Leu Asp Gly Gly Ala Pro Phe Tyr Thr Thr Tyr Arg
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Thr Ala Asp Gly Glu Phe Met Ala Val Gly Ala Ile Glu Pro Gln Phe
225 230 235 240
Tyr Glu Leu Leu Ile Lys Gly Leu Gly Leu Lys Ser Asp Glu Leu Pro
245 250 255
Asn Gln Met Ser Met Asp Asp Trp Pro Glu Met Lys Lys Lys Phe Ala
260 265 270
Asp Val Phe Ala Lys Lys Thr Lys Ala Glu Trp Cys Gln Ile Phe Asp
275 280 285
Gly Thr Asp Ala Cys Val Thr Pro Val Leu Thr Phe Glu Glu Val Val
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His His Asp His Asn Lys Glu Arg Gly Ser Phe Ile Thr Ser Glu Glu
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Gln Asp Val Ser Pro Arg Pro Ala Pro Leu Leu Leu Asn Thr Pro Ala

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325 330 335  
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<210> 109  
 <211> 1524  
 <212> DNA  
 <213> Homo sapien

<400> 109

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<210> 110  
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 <212> DNA  
 <213> Homo sapien

<400> 110

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&lt;210&gt; 111.

&lt;211&gt; 1289

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 111

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&lt;210&gt; 112

&lt;211&gt; 115

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 112

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Arg	Arg	Val	Phe	Tyr	Arg	Pro	Tyr	Leu	Gln	Ile	Phe	Gly	Gln	Ile	Pro
		65			70				75					80	
Gln	Glu	Asp	Met	Asp	Val	Ala	Leu	Met	Glu	His	Ser	Asn	Cys	Ser	Ser
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Glu	Pro	Gly	Phe	Trp	Ala	His	Pro	Pro	Gly	Ala	Gln	Ala	Gly	Thr	Cys
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Val	Ser	Gln	Tyr	Ala	Asn	Trp	Leu	Val	Val	Leu	Leu	Leu	Val	Ile	Phe
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Ala	Pro	Pro	Phe	Ile	Val	Ile	Ser	His	Leu	Arg	Leu	Leu	Leu	Arg	Gln
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 Trp Glu Ser Val His Lys Glu Asn Phe Leu Leu Ala Arg Ala Arg Asp  
 225                      230                      235                      240  
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 245                      250                      255  
 Asp Leu Ala Leu Lys Gln Leu Gly His Ile Arg Glu Tyr Glu Gln Arg  
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 Leu Lys Val Leu Glu Arg Glu Val Gln Gln Cys Ser Arg Val Leu Gly  
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<210> 113  
 <211> 553  
 <212> PRT  
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<400> 113  
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 20                      25                      30  
 Ala Ala Gly Ile Thr Tyr Val Pro Pro Leu Leu Leu Glu Val Gly Val  
 35                      40                      45  
 Glu Glu Lys Phe Met Thr Met Val Leu Gly Ile Gly Pro Val Leu Gly  
 50                      55                      60  
 Leu Val Cys Val Pro Leu Leu Gly Ser Ala Ser Asp His Trp Arg Gly  
 65                      70                      75                      80  
 Arg Tyr Gly Arg Arg Arg Pro Phe Ile Trp Ala Leu Ser Leu Gly Ile  
 85                      90                      95  
 Leu Leu Ser Leu Phe Leu Ile Pro Arg Ala Gly Trp Leu Ala Gly Leu  
 100                      105                      110  
 Leu Cys Pro Asp Pro Arg Pro Leu Glu Leu Ala Leu Leu Ile Leu Gly  
 115                      120                      125  
 Val Gly Leu Leu Asp Phe Cys Gly Gln Val Cys Phe Thr Pro Leu Glu  
 130                      135                      140  
 Ala Leu Leu Ser Asp Leu Phe Arg Asp Pro Asp His Cys Arg Gln Ala  
 145                      150                      155                      160  
 Tyr Ser Val Tyr Ala Phe Met Ile Ser Leu Gly Gly Cys Leu Gly Tyr  
 165                      170                      175  
 Leu Leu Pro Ala Ile Asp Trp Asp Thr Ser Ala Leu Ala Pro Tyr Leu  
 180                      185                      190  
 Gly Thr Gln Glu Glu Cys Leu Phe Gly Leu Leu Thr Leu Ile Phe Leu  
 195                      200                      205  
 Thr Cys Val Ala Ala Thr Leu Leu Val Ala Glu Glu Ala Ala Leu Gly  
 210                      215                      220  
 Pro Thr Glu Pro Ala Glu Gly Leu Ser Ala Pro Ser Leu Ser Pro His  
 225                      230                      235                      240  
 Cys Cys Pro Cys Arg Ala Arg Leu Ala Phe Arg Asn Leu Gly Ala Leu  
 245                      250                      255  
 Leu Pro Arg Leu His Gln Leu Cys Cys Arg Met Pro Arg Thr Leu Arg

260 265 270  
 Arg Leu Phe Val Ala Glu Leu Cys Ser Trp Met Ala Leu Met Thr Phe  
 275 280 285  
 Thr Leu Phe Tyr Thr Asp Phe Val Gly Glu Gly Leu Tyr Gln Gly Val  
 290 295 300  
 Pro Arg Ala Glu Pro Gly Thr Glu Ala Arg Arg His Tyr Asp Glu Gly  
 305 310 315 320  
 Val Arg Met Gly Ser Leu Gly Leu Phe Leu Gln Cys Ala Ile Ser Leu  
 325 330 335  
 Val Phe Ser Leu Val Met Asp Arg Leu Val Gln Arg Phe Gly Thr Arg  
 340 345 350  
 Ala Val Tyr Leu Ala Ser Val Ala Ala Phe Pro Val Ala Ala Gly Ala  
 355 360 365  
 Thr Cys Leu Ser His Ser Val Ala Val Val Thr Ala Ser Ala Ala Leu  
 370 375 380  
 Thr Gly Phe Thr Phe Ser Ala Leu Gln Ile Leu Pro Tyr Thr Leu Ala  
 385 390 395 400  
 Ser Leu Tyr His Arg Glu Lys Gln Val Phe Leu Pro Lys Tyr Arg Gly  
 405 410 415  
 Asp Thr Gly Gly Ala Ser Ser Glu Asp Ser Leu Met Thr Ser Phe Leu  
 420 425 430  
 Pro Gly Pro Lys Pro Gly Ala Pro Phe Pro Asn Gly His Val Gly Ala  
 435 440 445  
 Gly Gly Ser Gly Leu Leu Pro Pro Pro Ala Leu Cys Gly Ala Ser  
 450 455 460  
 Ala Cys Asp Val Ser Val Arg Val Val Val Gly Glu Pro Thr Glu Ala  
 465 470 475 480  
 Arg Val Val Pro Gly Arg Gly Ile Cys Leu Asp Leu Ala Ile Leu Asp  
 485 490 495  
 Ser Ala Phe Leu Leu Ser Gln Val Ala Pro Ser Leu Phe Met Gly Ser  
 500 505 510  
 Ile Val Gln Leu Ser Gln Ser Val Thr Ala Tyr Met Val Ser Ala Ala  
 515 520 525  
 Gly Leu Gly Leu Val Ala Ile Tyr Phe Ala Thr Gln Val Val Phe Asp  
 530 535 540  
 Lys Ser Asp Leu Ala Lys Tyr Ser Ala  
 545 550

&lt;210&gt; 114

&lt;211&gt; 241

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 114

Met Gln Cys Phe Ser Phe Ile Lys Thr Met Met Ile Leu Phe Asn Leu  
 1 5 10 15  
 Leu Ile Phe Leu Cys Gly Ala Ala Leu Leu Ala Val Gly Ile Trp Val  
 20 25 30  
 Ser Ile Asp Gly Ala Ser Phe Leu Lys Ile Phe Gly Pro Leu Ser Ser  
 35 40 45  
 Ser Ala Met Gln Phe Val Asn Val Gly Tyr Phe Leu Ile Ala Ala Gly  
 50 55 60  
 Val Val val Phe Ala Leu Gly Phe Leu Gly Cys Tyr Gly Ala Lys Thr  
 65 70 75 80  
 Glu Ser Lys Cys Ala Leu Val Thr Phe Phe Phe Ile Leu Leu Leu Ile



	85		90		95										
Phe	Ile	Ala	Glu	Val	Ala	Ala	Ala	Val	Val	Ala	Leu	Val	Tyr	Thr	Thr
	100							105					110		
Met	Ala	Glu	His	Phe	Leu	Thr	Leu	Leu	Val	Val	Pro	Ala	Ile	Lys	Lys
	115							120					125		
Asp	Tyr	Gly	Ser	Glu	Glu	Asp	Phe	Thr	Gln	Val	Trp	Asn	Thr	Thr	Met
	130							135					140		
Lys	Gly	Leu	Lys	Cys	Cys	Gly	Phe	Thr	Asn	Tyr	Thr	Asp	Phe	Glu	Asp
	145							150				155			160
Ser	Pro	Tyr	Phe	Lys	Glu	Asn	Ser	Ala	Phe	Pro	Pro	Phe	Cys	Cys	Asn
								165							175
Asp	Asn	Val	Thr	Asn	Thr	Ala	Asn	Glu	Thr	Cys	Thr	Lys	Gln	Lys	Ala
								180							190
His	Asp	Gln	Lys	Val	Glu	Gly	Cys	Phe	Asn	Gln	Leu	Leu	Tyr	Asp	Ile
	195							200					205		
Arg	Thr	Asn	Ala	Val	Thr	Val	Gly	Gly	Val	Ala	Ala	Gly	Ile	Gly	Gly
	210							215					220		
Leu	Glu	Leu	Ala	Ala	Met	Ile	Val	Ser	Met	Tyr	Leu	Tyr	Cys	Asn	Leu
	225							230					235		240
Gln															

<210> 115  
 <211> 366  
 <212> DNA  
 <213> Homo sapien

<400> 115  
 gctctttctc tccctctctc tgaatttaac tctttcaact tgcattttgc aaggattaca 60  
 catttcactg tcatgtatat tgtgttgcaa aaaaaaahaa gtgtctttgt ttaaaattac 120  
 ttggtttggt aatccatctt gctttttccc catttggact agtcattaac ccattctctga 180  
 actggtagaa aaacatctga agagctagtc taccagcacc tgacagggtg attggatagt 240  
 tctcagaacc atttcarcca gacagcctgt ttctatcttg tttaactaat tagtttgggt 300  
 tctctacatg cctaacaaac cctgtctcaa tctgtccat aaggtctgt gacttgaagt 360  
 ttagtc 366

<210> 116  
 <211> 282  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (282)  
 <223> n - A,T,C or G

<400> 116  
 acaagatga accatttctt atattataga aaaaataaaa tctaccctga ttctaatatt 60  
 gagaatgag atnaaacaca atttatata gtctacttag agaagatcaa gtgacutcaa 120  
 agattttact attttcatat tttaagacac atgattttct ctatttttagt aacctgggtc 180  
 atcgttttaa caaaggataa tgtgaacagc agagaggatt tgttggcaga aaatctatgt 240  
 tcaatctnga acLatctana tcacagacat tttatttctt tt 282

<210> 117  
 <211> 305

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (305)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 117

acacatgtcg	cttcaactgcc	tctcttagatg	cttctgtgtca	acatonagga	acagggacca	60
tatttatcc	ccctccctgaa	acaattgcag	aataanacaa	aattatctga	acaattgcag	120
ataaaggcaa	atatatatgaa	acacacaggtc	tgcagatatt	ggaaatcagt	caatgaagga	180
tactgatccc	tgatcaatgt	cctaattgcag	gatytyggaa	acagatgagg	tcacctctgt	240
gaatgcacca	gcttactgcc	tgtacagagt	ttctangctg	cagttcagac	agggagaaat	300
tggtt						305

&lt;210&gt; 118

&lt;211&gt; 71

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (71)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 118

accaaggtgt	atgaatctct	gaagtgggga	tctctgattc	cgcacaaac	tgagtggaaa	60
aantctctgg	t					71

&lt;210&gt; 119

&lt;211&gt; 212

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (212)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 119

actcgggttg	gtgtcagcag	gaagtggcat	tgaacatngc	aatgtggagc	ccaaaccaca	60
gaaaatgggg	tgaattggc	caactttcta	tnaacttatg	tggcaantt	tgcacacac	120
agtaagctgg	cccttcta	aaagaaaat	tgaaggttt	ctactaenc	ggattaant	180
aatggantca	aganactccc	aggcctcagc	gt			212

&lt;210&gt; 120

&lt;211&gt; 90

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (90)

&lt;223&gt; n = A,T,C or G

<400> 120  
 ectcgtttgca natcagggggc cccccagagt caccgttgcg ggagtccttc tggctttgcc 60  
 ctccgcgggc gcagaaatg ctgggggtgt 90

<210> 121  
 <211> 218  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{218}  
 <223> n = A,T,C or G

<400> 121  
 tgtcncgtga anacgacaga nagggttgtc aaaaatggag aanccttgaa gtcattttga 60  
 gaataagatt tgcataaaga ttgggggcta aaacatgggt attgggagac atttctgaag 120  
 atatncangt aaattangga atgaattcat ggttcttttg ggaattcctt taagatngcc 180  
 agcatanact tcatgtgggg atancagcta ccttctga 218

<210> 122  
 <211> 171  
 <212> DNA  
 <213> Homo sapien

<400> 122  
 taggggtgta tycactgta aggacaaaaa ttgagactca actggcttaa ccaataaagg 60  
 catttggttag ctcatgggac aggaagtcgg atgggtgggg atcttcagla ctgcatgagt 120  
 caccaccccg gcgggggtcat ctgtgccaca ggtccctgtt gacagtgcgg t 171

<210> 123  
 <211> 76  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{76}  
 <223> n = A,T,C or G

<400> 123  
 tctagcgtga agacnacaga atggtgtgtg ctgtgctatc caggaaacaa tttattatca 60  
 ttatcaanta ttgtgt 76

<210> 124  
 <211> 131  
 <212> DNA  
 <213> Homo sapien

<400> 124  
 acccttcccc aaggccaatg tctgtgtgtc taactggcgg gctgcaggac agctgcaatt 60  
 caatgtgtgt ggtcatatgg aggggaggag actctaaaat agccaatttt attctcttgg 120  
 ttaagatttg t 131

<210> 125  
 <211> 432  
 <212> DNA  
 <213> Homo sapien

<400> 125  
 acatttatcta ctggctatga aataqatggt ggaaaattgc gttaccaact ataccactgg 60  
 cttagaaaag aggtgatago tttcagagg acttgtgact ttgctcaga tgcagaaga 120  
 ctacagtcctg catctggrag aatguagat gaatttggat taatgagga tgcagaagat 180  
 ttgctcacc aaacaaagt gaaacaactg agagaaaatt ttcaggaaa aagacagtgg 240  
 ctcttgaggt atcagtcact tttagaagt ttcttagtt actgcatact tcatggatcc 300  
 catggtaggg gtcttgcatc tgaagaatg gaattgatt tgcctttgca agaattcag 360  
 caggaaacat cagaaccact atttctatga cctctgtcag agcaaacctc agtgccttc 420  
 ctctttgatt gt 432

<210> 126  
 <211> 112  
 <212> DNA  
 <213> Homo sapien

<400> 126  
 acacaacttg aatagtaaaa tgaactga gctgaattt ctaattcact ttctaaccat 60  
 agtaagaatg atatttcacc ccagggatca ccaatattt ataaaattt gt 112

<210> 127  
 <211> 54  
 <212> DNA  
 <213> Homo sapien

<400> 127  
 accacgaac cacaacaag atggaagcat caatccact gccagcaca gcag 54

<210> 128  
 <211> 323  
 <212> DNA  
 <213> Homo sapien

<400> 128  
 agctcattag taattgtttt gttgtttcat tttttctaa tgtctccct ctaccagctc 60  
 acctgagaca acagatgaa aatggaagg cagccagatt tctctttgc tctctgctca 120  
 ttctctctga agtctaggtt accattttt gggaccatt ataggcaata aacacagttc 180  
 ccacagcatt tggacagttt ctgtgttgt tttagaatg ttttctttt tcttagcctt 240  
 ttctgcaaa aggcctact agtcccttgc ttgtcagtg gactgggctc ccagggcct 300  
 aggcctgctt cttttccatg tcc 323

<210> 129  
 <211> 192  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{192}  
 <223> U = A,T,C or G

&lt;400&gt; 129

acatacatgtc gtgtatattt ttaaataatca utttgtatc actctgaatt tttagcatac	60
tgaataacata cttaacataat ttntgtgaac catgatcaga tacaacccaa atcatttate	120
tggcacatto atctgtgata naaagatagg tgaatttcat ttccttcaag ttggccaatg	180
gataaacaac gt	192

&lt;210&gt; 130

&lt;211&gt; 352

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(362)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 130

ccctttttta tgaatgagt agactgtatg ttctgaanatt tancacacac ctctctgaca	60
tataatgacg caacaaaaag gtgtgtgtta gtctatggt tcagtttatg cccctgacaa	120
gttctcattg tgttttgccg atcttctggc taatcgttgt atcttccatg ttattagtaa	180
ttctgtatc catcttgcta aggtctgga gatgtaacct gctangaggc taactttata	240
cttattttaa agctcttatt ttgttggtcat taaaatggca atttatgtgc agcactttat	300
tgcagcggga agcactgtg ggttgaattg aaagctctt gctaacttta aaaagtaatg	360
gg	362

&lt;210&gt; 131

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(332)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 131

ctttttgaaa gatcgtgtcc actcctgtgg acatcttgtt ttaatggagt ttcccatgca	60
gtangacttg tatggttgca gctgtccaga taaaaacatt tgaagagctc caaatgaga	120
gttctccrag gttcgcctg ctgtcccaag tctcagcagc agcctctttt agyaggcatc	180
ttctgaaata gattaaggca gcttgtaaat ctgatgtgat ttggtttatt atccaactaa	240
cttccatctg ttctcactgg agaaagccca gactcccan gacnggtacg gattgtgggc	300
atanaaggat tgggtgaagc tggcgtttgt gt	332

&lt;210&gt; 132

&lt;211&gt; 322

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(322)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 132

acttttgcca ttttgtatat ataaacaatc ttgggaatt ctctgaaaa ctgggtgtcc	60
---	----

agtggctaag	agaactcagat	ttcangcaat	tctgaaggga	aaaccagcat	gacacagaat.	120
ctcaaatctc	caaacagggg	ctctgtggga	aaactgaggg	aggacctttg	tatctcgggt	180
tttagcaagt	taaaatgaen	atgacaggaa	aggcttattt	atcaacaaag	agaagagttg	240
ggatgcttct	aaaaaaaact	ttggtagaga	aaatagggaat	gctnaatct	agggaagct	300
gtacacatct	acaattggtc	ca				322

&lt;210&gt; 133

&lt;211&gt; 278

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(278)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 133

acaagccttc	acaggtttta	ctaaattggg	attaatcttt	ctgtanttat	ctgcataatt	60
cttgtttttc	tttccatctg	gctcctgggt	tgacaatttg	tggaacaaac	tctattgcta	120
ctatttaaaa	aaatcacaa	atctttccct	ctaaagctatg	ctnaattcaa	actattcctg	180
ctatttctgt	tttgtcaag	aaattatatt	tttcaaaata	tgtntatttg	tttgatgggt	240
cccacgaac	actaataaa	accacagaga	ccagcctg			278

&lt;210&gt; 134

&lt;211&gt; 121

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(121)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 134

gtttanaaaa	cttggtttagc	tcctatagagg	aaagaatggt	aaactttgta	ttttaaaaa	60
tgattctctg	aggttaaaact	tggttttcaa	atgttatttt	laattgtatt	ttgcltttgg	120
t						121

&lt;210&gt; 135

&lt;211&gt; 350

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(350)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 135

acttanaacc	atgcctagca	catcagaatc	cttcaagaa	catcagttata	atnclatacc	60
atancaagtg	gtgactgggt	aaagctgcga	caaaggtcag	ctggcacatt	acttgtgtgc	120
aaacttgata	cttttgttct	aagttagaac	tagtatacag	tnootaggan	tgglactcca	180
gggtgcccc	aaactcctgc	agccgtctct	ctgtgccagn	ccrtgnaagg	aactttcgt	240
ccacctcaat	caagccctgg	gccatgctac	ctgcaattgg	ctgaacaaac	gttlgctgag	300
ttcccaagga	tgcaagcct	ggtgctcaac	tcttggggcg	tcaactcagt		350

<210> 136  
 <211> 399  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(399)  
 <223> n = A,T,C or G

<400> 136  
 tgtacggtga agacgacaga agttgcatgg cagggacagg gcaggggcga ggccagggtt 60  
 gctgtgattg tatccgaata ntctctgtga gaaaagatga tgagatgacg tgagcagcct 120  
 gcagacttgt gtctgcttc aanaagccag acagggaaggc cctgcttgc ttaggtctga 180  
 cctggcgggc agccagccag ccacagggtg gcttcttct tttgtggtga caaccccag 240  
 aaaactgcag agggccaggg tcagggtatg gtgggtangl gccataaaa cccagggtgc 300  
 tccagggaac ccgggcagag gccatccca cctccagcca gcctgcccac tggcgtgatg 360  
 ggtgcagang gatgaagcag ccagntgctc tctctgtgt 399

<210> 137  
 <211> 165  
 <212> DNA  
 <213> Homo sapien  
 <220>  
 <221> misc\_feature  
 <222> (1)...(165)  
 <223> n = A,T,C or G

<400> 137  
 actggtgtgg tngggaggtg tctgtgtgtt acaagttgan gtgacttcac gatggtgtgt 60  
 ggaggaggtg tctgaacgta gggatgtaga ngttttggcc gtgctaaatg agcttoggga 120  
 ttggtgtgtc ccactggttg tcaatgtcat tgggtgggtt cctgt. 165

<210> 138  
 <211> 338  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(338)  
 <223> n = A,T,C or G

<400> 138  
 actcactgga atgccacatt cacaacagaa tcagagggtc gtagaaacat taatggctcc 60  
 ttaacttctc cagtaagaat cagggacttg aatggaaac gttacagcc acatgccraa 120  
 tcttgggcag tctccatgc ctccacagt gaaagggtt gaggaaatc acatccaatg 180  
 tcatgtgttt ccagccacac caaaggtgc ttgggttga gggctgggg catananggt 240  
 cagcctcag gaagcctcaa gttccattca gctttgccac tgtacattcc ccatnttaa 300  
 aaaaactgat gccctttttt ttttttttg taaaattc 338

<210> 139  
 <211> 382

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 133

gggaatcttg	gtttttggca	tctggtttgc	ctctagccga	ggccactttg	acagaacaaa	60
gaaggaggact	tcgagtaaga	aggagattta	cagccagccl	agtgcacgaa	gtggaaggaga	120
attcaccacag	acctcgctcat	tcctgggtgfg	aggctggctg	gtcaccgcgc	tatcatctgr	180
atttgccctta	ctcaggtgct	accggactct	ggcccttgat	gtctgtagtt	tcacaggatg	240
acttatcttgc	cttctacac	ccacaggggc	ccctacttcc	tcggatgcat	ttttaataat	300
gtcagctatg	tgcacacac	tccttcctgc	ctcctctcc	tttctacaa	ctgctgagtg	360
gcctggaaact	tgtttaaagt	gt				382

&lt;210&gt; 140

&lt;211&gt; 200

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (200)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 140

accaaactt	ctttctgtt	tgctngattc	taatataggg	gtctngcttn	ttctaaanat	60
acttttcatt	taacancttt	tgttaagtgt	caggttgca	tttgcctcat	anaattattg	120
ttttacacct	tcacattgta	tgtgtttgtc	tccttanaga	ttggtgaaat	cacatatttt	180
atattcagca	taaaggagaa					200

&lt;210&gt; 141

&lt;211&gt; 335

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (335)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 141

actttatttt	caaacacctc	atatgttgca	aaaacacat	agaaaaataa	agtttggttg	60
gggtgctgac	tzaacttcaa	gtcacagact	tttatgtgar	agattggagc	agggtttglt	120
atgratgtag	agaacccaaa	ctaatttatt	aaacaggata	gaacaggct	gtctgggtga	180
aatggttctg	agaacratcc	aattcacctg	tcagatgctg	atanactagc	tccttcagatg	240
tttttctacc	agttcagaga	tinggttaatg	actantcca	atgggggaaa	agcaagatgg	300
attcacaaar	caagtaattt	laaacaaaga	cactt			335

&lt;210&gt; 142

&lt;211&gt; 459

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (459)

&lt;223&gt; n = A, T, C or G



&lt;400&gt; 142

accaggthaa	lattgocaa	tatatcttt	cnaattggg	gctaaacaga	vgtgattta	60
gggttgttta	aagacaaccc	agcttaatat	cagagaaat	tgtgaccttt	catggagtat	120
ctgatggaga	aaacactgag	tttgacaaa	tcttattta	ttcagatagc	agtctgatca	180
caatyggtcc	aacaacactc	aaataataa	tcaatataa	tcagatgta	aagattggtc	240
ttcaaacatc	atagccaatg	atgcccgcct	tgcctataat	ctctccgaca	taaaaccaca	300
tcaacacctc	agtggccacc	aaacattca	gcacagcttc	cttaactgtg	agctgtttga	360
agctaccagt	ctgagcaata	ttgactatnt	ttttcangct	ctgaatagct	ctagggatcl	420
cagcangggg	gggaggaacc	agctcaacct	tggegtant			459

&lt;210&gt; 143

&lt;211&gt; 140

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 143

acatttcctt	caaccaagtc	aggactctctg	gcttctgtgg	gagttcttat	caactgaggg	60
aatccaaac	agtctctct	agaaaggaat	agtgtcaaca	accacacaa	ctctccctgag	120
accatccgac	ttccctgtgt					140

&lt;210&gt; 144

&lt;211&gt; 164

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (164)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 144

acttcagtaa	caacatataa	taacaacatt	aaagtgttat	tgcctctttt	gtcattttct	60
atctatacca	ctctcccttc	tgaacaaan	aatcactano	caatcactta	tacaaatttg	120
aggaatttaa	tcaatatttg	ttttcaataa	ggaaaang	atgt		164

&lt;210&gt; 145

&lt;211&gt; 303

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (303)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 145

acgtagacca	tccaactttg	tatttgtaat	ggcaaacatc	cagnagcaat	tctaaacaa	60
actggagggt	atttatccc	aattatccc	ttcattcaaa	tgcctctctc	ctcaggctat	120
gcaggacagc	tatcataagt	gggcccaggg	atccagatcc	ttccatttgt	ataaacctta	180
gtaggggagt	ccatccaaat	gacaggtcta	atcnaaggag	gaatgggac	ataagrcag	240
cagtataatn	ttgcttagct	gaacacagca	caaaagactt	accgcccgtg	tgattaccat	300
caa						303

&lt;210&gt; 146

&lt;211&gt; 327

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(327)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 146

actgcagctc aattagaagt ggtctctgac ttctatcanc ttctccctgg gttccatgac	60
actggccctgg agtgactcat tgctctgggt gggtgagaga gtccctttgc caacaggcct	120
craagtccgg gctgggattt gtttcttthc cacattctag caacaatag ctggccactt	180
cctgaacagg gagggtggga ggagccagca tggacaaggc tgcactttc taaagtagec	240
agacttgccc ctgggacctg cacacctact gatgaacttc tgtgacctga ggatgggaatg	300
tggggctgag ctgtgtgact ctatgggt	327

&lt;210&gt; 147

&lt;211&gt; 173

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(173)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 147

acattgtttt tttagatatc agcattgena gagctctcct taavgtgaca caatgggaag	60
actgggaacc ataccacat ctctgttctg agggataatt ttctgataaa gtctttgtgt	120
atattcaagc acatatgta tatattatc agttccatgt ttatagcta gtt	173

&lt;210&gt; 148

&lt;211&gt; 477

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(477)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 148

acaaacctt tatctcatcg aatttttaac ccaaactcac tcaactgtgc ttctatctct	60
atgggaatata ttatttgatg ctccatttca tcaacatat atgaataata cactcatact	120
gccctactac ctgctgcact aatcacattc ccttctgtc ctgacctga agcattggg	180
gtggtcttag tggccatcag tccanycctg cacttgagc ccttgagctc cattgctcac	240
acacacccac ctacacgacc caatcctctt acacagctac ctcttggctc tctaacccca	300
tggattatnt ccaatttcag tcaattaaat tectttaac actctaccuy acatgtccag	360
caccactggt aagccttctc cagccaacac acacacacac acacacacac ccacacatat	420
ccaggcacag gctacctcat ctccacaatc acccctttaa ttacccatgct atgggtgg	477

&lt;210&gt; 149

&lt;211&gt; 207

&lt;212&gt; DNA

<213> Homo sapien

<400> 149

```
acagttgtat tataatctca agaatataar ttgcantgag agcatttaay agggagaagac      60
taacgtatatt tagagagcca aggaaggttt ctgtggggag tgggatgtaa ggtggggcct      120
gatgataaat aagagtcagc caggttaagt ggtggtgtgt tatgggcaca gtgaagaaca      180
tttcagggag agggacacag agtgaa
```

207

<210> 150

<211> 111

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> {1}...{111}

<223> n = A,T,C or G

<400> 150

```
accttgattt cattgctgct ctgatggaaa ccccaactat taatttagct aagacatggg      60
cacttaabtg tggcagtggt ttggacttgt taactantgg catctttggg t
```

111

<210> 151

<211> 196

<212> DNA

<213> Homo sapien

<400> 151

```
agcgcgag gtcatttga acattccaga taactatcat tactcgatgc tgttgataac      60
agcaagatgg ctctgaactc aggggcacaa caagctattg gcccttacta tgaaaacat      120
ggataccaa cggaaaacc ctatccgca cagcccactg tggccccac tttctacag      180
gtgcatcgg ctcagt
```

196

<210> 152

<211> 132

<212> DNA

<213> Homo sapien

<400> 152

```
acagcattt cacatgtaag aaggagaaa ttcttaaatg tagggagag ataacagac      60
cttccattt tcatctagt gtggaaacct gatgctttat gttgacagga atgaaccag      120
gaggagctt gt
```

132

<210> 153

<211> 285

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> {1}...{285}

<223> n = A,T,C or G

<400> 153

```
acaaaccca nganaggcca ctggccgtgg tgcctggcc tccaaacatg aaagtgtcag      60
```

cttctgctct tatgtctca tctgcaact cttaccatt ttatctctg ctcaagcagga	120
gcacatcaat aaagtccaaa gtcttggaact tggccttggc ttggaggag ttatcaacac	180
cctggctagt gaggggtggg cggcgctcct ggaatgacgg atctgtgaag tctgtcacca	240
gtctgcaggc cctgtggagg cgccgtccac agggagtnag gaatt	285

<210> 154  
 <211> 333  
 <212> DNA  
 <213> Homo sapien

<400> 154	
accacagtcn tgbtgggcca gggcttcatt acccctctctg tgaaaagcca tattatcaac	60
accccaaat tttcctttaa catccttaac tgaaggggtc agcctcttga ctgcacagac	120
cctaagcggg ttacacagct aactccact ggccttgatt tgtgaaattg ctgtgctctg	180
attggacag gactcgaagg tgttcagctc cctctctcag tggacagaga ctctgatttg	240
agtttcacaa attctcgggc cactctgta tctctctct gaaataaat ccggagaatg	300
gtcaggcctg tctatccat atggatcttc cgg	333

<210> 155  
 <211> 308  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> 11... (308)  
 <223> n = A,T,C or G

<400> 155	
actggcaata ataaaaccca catcacagtg ttgtgtcaca gatcatcagg gcatggatgg	60
gaaagtgett tgggaactgt aaagtgccta acacatgac gatgattttt gttataatat	120
ttgaatcagg gtgcatacaa actctctctg ctgtctctcc tgggccccag cccacagccc	180
atcacagctc actgtctgt tcatccaggg ccagcatgta gtggctgatt ctctctgggt	240
gtttctagcc tccanaagtt tctctgaagg caacccaaac tctangtata aggcattgctg	300
ggcctggg	308

<210> 156  
 <211> 295  
 <212> DNA  
 <213> Homo sapien

<400> 156	
acottgctcg gtgcttggaa catattagga actcaaaata tggatgata acagtgccta	60
ttcttgatta ctgagagaac tgttagacat ttagtgaag attttctaca caggaaactga	120
gaataggaga ttatgtttgg cctcataat ctctctatc ctcttgcct cattctatgt	180
ctaatatatt ctcaatcaa taaggctagc ataacagga aatcgacaa ataccaatat	240
aaaaccagat gtctatctct aagattttca catagaaac aaattacag actat	295

<210> 157  
 <211> 126  
 <212> DNA  
 <213> Homo sapien

<400> 157	
ccaagtttaa atagtgtgt cactgtgcat gtgctgaat gtgaaatcca ccacattct	60

gaagagcaaa acaaattctg tcatgtaata totatcttgg gtcatggga tatctgtccc 120  
cttagt 126

<210> 158  
<211> 442  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> {1}... (442)  
<223> n = A,T,C or G

<400> 158  
accactggt cttggaaaca cccatcctta atacgatgat ttttctgtcg tctgaaatg 60  
aanccagcag gctgcuccata gtcagtccct ccttccagag aaaaagagat ttgagaaagt 120  
gctggggtta ttcaccatta atttcctccc ccaactctc tgaattttcc cttactattt 180  
ctggtaggttc tgacaaaagg aggtcatggt ttgttgagca ttgggatcc cagtgaagta 240  
natgtttgta gctttgcata cttagccctt cccacgcaca aacggagtg cagagtggtg 300  
ccaacctgt ttcccaagt cactagaca gattcacagt ggggaattct ggaagctgga 360  
nacagagggg ctctttgcag agccgggact ctgagangga catgagggcc tctgctctg 420  
tgttcattct ctgatgtcct gt 442

<210> 159  
<211> 498  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> {1}... (498)  
<223> n = A,T,C or G

<400> 159  
acttccaggt aacgtgtgtg ttctcgttga gctgaactg atgggtgacg ttgtaggttc 60  
tccaaacaaga actgaggttg cagagcgggt agggaaagat gctgttccag ttgcacctgg 120  
gctgctgtgg actgtgtgtg attctcact acggcaccag gttgtggaac tggcannaag 180  
gtgtgtgttt gganttgagc tggggcggct gtggtaggtt gtgggtcttt caacaggggc 240  
tgetgtgtgt cggggangtg aanygtgtgt gtracttgag cttggccagc tctggaagt 300  
antanattct tctgaagge cagcgttgt ggaagctggca agggtcantg ttgtgtgtaa 360  
cgaaccagtg ctgctatggg tgggtgtana tcttccacaa agcctgaagt tatggtgtcn 420  
tcaggtaana atgctggttc agtgcctctg ggcnctgtg gaagggtgtg nattgtcacc 480  
aagggaataa gctgtgtg 498

<210> 160  
<211> 380  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> {1}... (380)  
<223> n = A,T,C or G

<400> 160

```

accctgcaccc agcttccctg ccaaaactaac aaggagacat caacctctag acagggaaac      60
agcttcagga tacttccagg agacagagcc accagcagca aaacaaatat tccuatgctt      120
ggagcatggc atagaggaag ctganaaetg tggggtctga ggaagccatt tgagtctagg      180
cactngacat ctcatcagcc acttggtgtga agagatgccc catgacccca gatgcctctc      240
cagaccttac ctccatctca cacacttgag ctctccartc tgtataattc taacatccctg      300
gagaaaaatg gragtttgac cgaacctgtt cacaacgcta gaggetgatt tctaacgaaa      360
cttgtagaal gaagcctgga                                     380

```

<210> 161  
 <211> 114  
 <212> DNA  
 <213> Homo sapien

```

<400> 161
actccacatc cctctgagc agggggcttg cgttcaaggc gtatttgccc ttgcccgtca      60
cactgtccac tggccrctta tccatttggc gcttaattcc tcgaaagagc atgt      114

```

<210> 162  
 <211> 177  
 <212> DNA  
 <213> Homo sapien

```

<400> 162
accttctgaa tcgaaacaaa tgatacttag ttagtcttta atatcctcat atatctcaaa      60
gttttactac tctgataatt ttgtaaacca ggtaaccaga acatccagtc atacagcttt      120
tggtgatata taacttggca ataaccagc ctggtgatcc ataaaactac tcaactgt      177

```

<210> 163  
 <211> 137  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{137}  
 <223> n = A,T,C or G

```

<400> 163
catttatada gacaggcgtg aagacattca cgacaaaaac gggaaattct atcccgtgac      60
canagaaggc agctacggct actctacat cctggcgctg gtggccttcg catgcacctt      120
catcagcggc atgatgt                                     137

```

<210> 164  
 <211> 469  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> {1}...{469}  
 <223> n = A,T,C or G

```

<400> 164
cttatcaca tgaaatttct cctgggagcc gttgtgatct ttgccacatt cgtgacttta      60
tgcaatgcac catgctatct catcctaat gaggagattc caggagattc aacaggaaa      120

```

```

tgcattggatc tcacagggaac caaacaccca ataaactcgg agtggcagac tgacaactgl      180
gagacatgca ctctgtacga aacagaaatt tcattgttgc ccttgtttc tacacctgtg      240
ggttatgaca aagacaactg ccaagaatc ltcaggagg aggaclgcaa gtatatctg      300
gtggagaaga aggaacacaa aaagacctgt tctgtcagtg aatggataat ctactgtgct      360
tctagttagg aagagggtcc caggccaggc ctactctcc tctggcctct aatagtcaat      420
gattgtgtag ccatgcttat cagtaaaag atctttgagc aaacacttt      469

```

&lt;210&gt; 165

&lt;211&gt; 195

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{195}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 165

```

acagtttttt atatatatcg acattgcagg cacttgtgtt cagtttcatz aagctgggtg      60
atcagctgtc atcactatt ccttggctag agtaaaaatt attcttatag cccatgtccc      120
tgcaggccgc ccgcccgtag ctctgtttcc agtcgtcttg gracacaggg tgcaggact      180
tcccttgaga tgagt      195

```

&lt;210&gt; 166

&lt;211&gt; 383

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{383}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 166

```

acactttagt agtgtggcac atcagggggc catcaggggc acagtcactc atagcctcgc      60
cgaggtcgga gtccacacca ccggtgtagg tgtgtcgaat cttgggcttg ggcgccacct      120
ttggagaagg gatatgctgc acacacatgt ccacaaagcc tgtgaactcg ccaagaatt      180
tttgcagacc agcctgaaga aggggtggat gttcagcttc agtcctctct tctcaggtg      240
gatgcuaacc tegtctangg tccgtgggaa gctgggtgter acntcaccta caacctgggc      300
gangatctta taaagaggct ccnagataaa ctccagaaa cttctctggg agctgctagt      360
nggggccttc ttggtgaact ttc      383

```

&lt;210&gt; 167

&lt;211&gt; 247

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{247}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 167

```

acagagccag accttggaca taatgaanc agagatteag actaaacccc aagtoganat      60
tggagcagaa actggagcaa gaagtgggcu tggggctgaa glagagacca aggcactgc      120

```

tatanccata caccagagcca accttcaggc caaggcctatg gttggggcag anccagagac	180
tcaatctgan tccaaagtgg tggctygaac actggctatg acanaggcag tgactctgac	240
tgangtc	247

&lt;210&gt; 168

&lt;211&gt; 273

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{273}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 168

acttctaagt ttctagaag tgggaagatt gtatcctcc tgaatttggg ttacttcaa	60
aatccctcan ccttggtctt cactactgtc tctactgana gtgtcatgtt tccacaaagg	120
gctgaacoot gagcttgnat ttctactcat ccttgagaag ccttttccag tagggtyggc	180
aattcccaac ttcttgcga caagcttccc aggtctcttc ccttggaacc ctccagcttg	240
agtcacagat aaactcatgg gtgcctctg gca	273

&lt;210&gt; 169

&lt;211&gt; 431

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{431}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 169

acagccttgg cttecccaaa ctccacagtc tcaagtgcga aagatcatct tccagcagtc	60
agctcagacc agggctcaag gatgtgacat caacagtttc tgytttcaga acaggttota	120
ctactgtcaa atgacccccc atacttcttc aaaggtctgt gtatgttttg cacaggtgag	180
ggcagcagaa agggggtant tactgatgga caccatcttc tctgtatart cccactgac	240
cttgccatgg gcaaaaggcc ctaccacaaa aacaatagga tcaatgcttg gcaccagctc	300
acgcacatca ctgacaaccg ggatygaaaa agaantgcca acttctatar atccaaactgg	360
aaagtgatct gatctggat tcttaattac ctccaaaagg ttchgggggg catcagctgc	420
tgaacactg a	431

&lt;210&gt; 170

&lt;211&gt; 266

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{266}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 170

acctgtgggc tgggtgttta tgctgtgcu ggtgtctgaa agggagttca gaggtggagc	60
tcaaggagct ctgcaggcat ttgccaanc ctctccanag canaggagc aacctaacct	120
cccgctaga aaagcaccag attggagtc tgggagggg agttggggtg ggcatttgat	180



gtatacttgt cacttgaatg aangagccng agaggaanga gacgaaatg analtygctt 210  
tcaaagctag gggctctgga gytgga 256

<210> 171

<211> 1248

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> {1}...{1248}

<223> n - A,T,C or G

<400> 171

```

ggcagccaaa tcataaacgg cgaggactgc agcccgact cguagccctg gcaaggcgga 60
ctggtcattg aaaaacgaatt gttctgctcg ggcgtccctg tgcattccga gtgggtgctg 120
ttagccgcac actgtttcca gaagtgaatg cagagctcct acaccatcgg gctgggcttg 180
cacagtcttg aggcgcacca agagccagg agccagatgg tggaggcccg cctctccgta 240
cgscacccag aytacacag acccttgctc gctacgacc tcatgtcat caagttggac 300
gaatccgtgt ccgagctctg caccatccgg agcatcagca ttgcttcga gtgcctacc 360
gcgggggaact ctgctcctgt ttctggctgg ggtctgctgg rgaacggcag aatgcctacc 420
gtgttgcagt gcgtgaacgt gtccgtggtg tctgaggagg tctgcagtaa gcttatgac 480
ccgtgttacc accccagcat gttctgcgac ggcggagggc aagaccagaa ggactcctgc 540
aacagtgaat ctgggggggc cctgatctgc aacgggtact tgcagggctt tgtgttttc 600
ggaaaagccc cgtgtgaca agttggctg ccaggtgtct acacnaacct ctgcaaatc 660
actgagtgga tagagaaaac cgtccaggcc agttaartc ggggactggg aacccatgaa 720
attgaccccc aatatcatcc tgcgggaagg attccaggaat atctgttccc agccctcct 780
ccctcaggcc caggagtcca ggcgcccgcc cctcctccc tcaaaccaag ggtacagatc 840
ccagccctct cctcctcag acuuaggagt ccagaccccc cagccctccc tccctcagac 900
ccaggagtcc aguccctct cctcagacc caggagtcca gaccccccag cccctcctcc 960
ctcagaccca ggggtccagg cccccaaccc cctcctccc agactragag gtccaagccc 1020
ccaaaccttc attcccca cccagaggtc cagggtccag cccctcctcc cctragacca 1080
gcagtccat gccacctaga ctntcctgt acacagtcc ccttgtggc acgttgcccc 1140
aaccttacca gttggtttt ctttttngt ccttttccc tagatccaga aataaagttt 1200
aaggagagng caaaaaaaa aaaaaaaa aaaaaaaa aaaaaaaa 1248

```

<210> 172

<211> 159

<212> PRT

<213> Homo sapien

<220>

<221> VARIANT

<222> {1}...{159}

<223> Xaa - Any Amino Acid

<400> 172

```

Met Val Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro
1          5          10          15
Leu Leu Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser
20          25          30
Glu Ser Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr
35          40          45
Ala Gly Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Gly
50          55          60

```

Arg Met Pro Thr Val Leu Gln Cys Val Asn Val Ser Val Val Ser Glu  
 65 70 75 80  
 Glu Val Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe  
 85 90 95  
 Cys Ala Gly Gly Gln Xaa Gln Xaa Asp Ser Cys Asn Gly Asp Ser  
 100 105 110  
 Gly Gly Pro Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe  
 115 120 125  
 Gly Lys Ala Pro Cys Gly Gln Val Gly Val Pro Gly Val Tyr Thr Asn  
 130 135 140  
 Leu Cys Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Ala Ser  
 145 150 155

<210> 173  
 <211> 1265  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)... (1265)  
 <223> n - A, T, C or G

<400> 173

ggcagccccc	actgcagccc	ctggcaggcc	gactgtgtca	tggaaaaaga	attgttctgc	60
tggggctcc	tgggtgcatc	gcagtgggtg	ctgtcagccc	ccactgtttt	ccagaactcc	120
tacaccatcg	ggctgggccc	gcacagtctt	gagggccgac	agagagccag	gagccagatg	180
gtggaggcca	gcctctccgt	acgggaaccca	gagtacaaca	gaccccttgc	cgctaacyac	240
ctratgtctc	tcaagtttga	cgatcccggt	tccgagctcg	acaccatccg	gagcatcagc	300
attgcttccg	agtgccttac	cggggggaaac	tattgctccg	cttctggctg	gggtctgctg	360
gcgaacgggt	agttccaggg	tgtgtgtctg	ccctcttcaa	ggaggccctc	tgcaccagtc	420
cgggggctga	ccagagatcc	tgggtcccag	gcagaatgcc	taccgtgctg	cagtgcgtga	480
acgtgtccgt	ggtgtctgag	gaggtctgca	gtaagctctc	tgaccgctcg	taccaaccca	540
gcctgttctg	cgccggcgga	gggcaagacc	agaaggactc	ctgcacaggt	gactctgggg	600
ggcccttgat	ctgcaacggg	tacttgccag	gccttgtgtc	tttcggaaaa	gcccctgtgt	660
gccaagttgg	ctgtccaggt	gtctacacca	acctctgcaa	attcactgag	tggatagaga	720
aaacgttcca	ggccagttca	ctctggggac	tgggaaccca	tgaattgac	ccccaaatcc	780
atcttgccga	aggaattcag	gaatatctgt	tcccagcccc	tctccctcca	ggccnaggag	840
tccagggccc	cagccctccc	tccctcaaac	caagggtaca	gatccccagc	ccctctctcc	900
tacagaccag	gagtcagagc	ccccagcccc	ctctctctcc	agaccagaga	gtccagcccc	960
tctctctcca	gacccaggag	tccagacccc	ccagccctcc	ctccctcaga	ccaggggggt	1020
gagggcccca	acccctctcc	cttcagagtc	agaggtccaa	gcccccaacc	ctctgttccc	1080
cagaccaga	ggttnaggtc	ccagccctcc	ttccctcaga	cccagnggtc	caatgcccac	1140
cagattttcc	ctgnacacag	tgcctccctg	tggngngttg	acccacccct	accagttggg	1200
ttttcatttt	tngtcccttt	cccttagatc	cagaaataaa	gtttcagaga	ngngcaaaaa	1260
aaaaa						1265

<210> 174  
 <211> 1459  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)... (1459)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 174

ggtcagccgc	acactgtttc	caggaagttag	tgcagagctc	ctacaccate	gggctgggac	60
tgcacagtct	tgagggcgac	cagagagcag	ggagccagat	ggtagaggcc	agcctctccg	120
taaggacccc	agagtacaac	agacccttgc	tcgtatacga	ccctcatgctc	atcaagtttg	180
acgaatccgt	gtccgagctc	gacaccatcc	ggagcatcag	cattgcttcg	cagtgcctta	240
cgcgggggaa	ctcttgcttc	gtttctggct	ggggtctgct	ggcgaaacgg	gagctccacg	300
gtgtgtgtct	gcccctcttc	aggaggtcct	ctgcccaytc	gcgggggctg	acccagagct	360
ctgcgtcccc	ggcagaatgc	ctaccgtgct	gcagtgcgtg	aaagtgtcgg	tggtgtctga	420
ngaggtctgc	antaagctct	atgaccgct	gtaccacccc	ancatgttct	gcgcggggcg	480
agggcaagac	cagaaggact	cctgcaacgt	ggagaggggg	aaaggggggg	gcaggcgact	540
cagggaaggg	tggaagaggg	ggagacagag	acacacaggg	ccgcacggcg	agatgcagag	600
atggagagac	acacagggag	acagtgaaca	ctagagagag	aaatgcagag	aaacagagaa	660
ataaacacag	gaataaagag	aagcaaagga	agagaagaa	agaaacagac	atggggaggc	720
agaaacacac	acacatagaa	atgcagttga	ccttccaaac	gcattggggc	tgagggtggg	780
gacctccacc	caataagaaa	tctctttata	acttttgact	ccccaaaaac	ctgactagaa	840
atagcctact	gttgacgggg	agccttacc	ataacataaa	taqtcgattt	atgcatacgt	900
tttatgcatt	catgatatac	ctttgttggg	attttttgat	atttctaacg	taacagttc	960
gtctgtgaat	tttttttaaa	tgttgcaact	ctcctaaat	ttttctgatg	tgtttcttga	1020
aaaaatccaa	gtataagtg	acttctgcat	tcaaacaggg	gttgttcaag	ggtcaactgt	1080
gtacccagag	ggaaacagtg	acacagatlc	atagaggtga	aacacgaaag	gaaacaggaa	1140
aaatcaagac	tctacaaaga	ggctgggcag	ggtggctcat	gcctgtaate	ccagcacttt	1200
gggaggcgag	gcagggcagat	cacttgaggt	agggagttca	agaccagcct	ggccaaaatg	1260
gtgaaatcct	gtctgtacta	aaaataaaaa	agttagctgg	atatygtggc	agggcgcctg	1320
aatccagct	acttgggagg	ctgaggcagg	agaattgctt	gaatatggga	ggcagaggtt	1380
gaagtgaagt	gagatcacac	cactatcttc	cagctggggc	aacagaytaa	gactctgtct	1440
caaaaaaaa	aaaaaaaaa					1459

&lt;210&gt; 175

&lt;211&gt; 1167

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(1167)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 175

ggcagccct	ggcagggcgc	actggctcatg	gaaaaagaat	tgttctgctc	gggctgctg	60
gtgcacccgc	agtgggtgct	gtcagccgca	cactgtttcc	agaaatccta	caccatcggt	120
ctgggcttgc	acagtcttga	ggcggaccaa	yagccagggg	gccagatggt	ggagggcagc	180
ctctccgtac	ggcaccacga	gtaccacaga	ctcttgcctg	ctaacgacct	catgctcate	240
aaagtggagc	aatccgtgtc	agagctctgac	accatcggga	gcatacgcat	tgcttcgacg	300
tgcctacccg	gggggaactc	ttgcctcgtn	tctggctggg	gtctgctggc	gaacggcaga	360
atgcctarcc	tgttgcactg	cgtgaacgtg	tgggtgggtg	ctgaggangt	ctgcagtaag	420
ctctatgac	cgtgttarca	ccccagcatg	ttctgcgcgc	gcggagggga	agacnagaag	480
gactcctgca	acgggtgaact	tggggggccc	ctgatctgca	acgggtactt	gcagggcctt	540
gtgtcttctg	gaaaagccrc	gtgtggccaa	cttggcgtgc	caggtgtcta	cacraacctc	600
tgcacatcca	ctgagtggt	agagaaaacc	gtccayncca	gttaactctg	gggactggga	660
acccatgaaa	ttgaccccc	aatacatcct	gcggaangaa	ttcagggaata	tctgttccca	720
gccctcctc	cttcaggccc	aggagtcag	gccccuagcc	cctctcccl	caaaccaagg	780
gtacagatcc	ccagccctc	ctccctcaga	cccaggagtc	cagacccccc	agccctcct	840
cctcagacc	caggagtcca	gccctctctc	cctcagagcc	aggagtcccg	accccccagc	900

```

cccccccccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc 960
cccccccccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc 1020
cccccccccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc 1080
cccccccccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc 1140
cccccccccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc ccccccaccc 1167

```

<210> 176  
 <211> 205  
 <212> PRT  
 <213> Homo sapien

<220>  
 <221> VARIANT  
 <222> (1)...[205]  
 <223> Xaa - Any Amino Acid

<400> 176  
 Met Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln Trp  
 1 5 10 15  
 Val Leu Ser Ala Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu  
 20 25 30  
 Gly Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val  
 35 40 45  
 Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Leu Leu Leu  
 50 55 60  
 Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu Ser  
 65 70 75 80  
 Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala Gly  
 85 90 95  
 Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Gly Arg Met  
 100 105 110  
 Pro Thr Val Leu His Cys Val Asn Val Ser Val Val Ser Glu Xaa Val  
 115 120 125  
 Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe Cys Ala  
 130 135 140  
 Gly Gly Gly Gln Asp Gln Lys Asp Ser Cys Asn Gly Asp Ser Gly Gly  
 145 150 155 160  
 Pro Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe Gly Lys  
 165 170 175  
 Ala Pro Cys Gly Gln Leu Gly Val Pro Gly Val Tyr Thr Asn Leu Cys  
 180 185 190  
 Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Xaa Ser  
 195 200 205

<210> 177  
 <211> 1119  
 <212> DNA  
 <213> Homo sapien

<400> 177  
 gggcactcgc aggcctggca gggcgcactg gtcctcggaa acgaattggt ctgctcgggc 60  
 gtcctcgggc atcgcagctg ggcctcgtca ggcgcacact gtctcagaa ctctacacac 120  
 atcgggcttg ggcctcagcg tcttgaggcc gcccaagagc caggagacga gatgctggag 180  
 gccagcctct ccgtacggca cccagagtaa aacagacacct tgcctcgtca cgcctcagc 240  
 ctcatcaagt tggacgacac cgtgctcagc tctgacacaa tccggagcat cagccttgc 300

```

tcgcagtgcc ctaccgaggg gaactcttgc ctggtttctg gctgggggtct gctggggaac 360
gatgctgtga ttgccatcca gtcccagact gtgggaggct gggagtgtga gaagctttcc 420
caaccctggc aggggttgtac catctcggca acctccagtg caaggagcgc ctgctgcac 480
ctcactgggt gctcactart gctcactgca tcaccgggaa cactgtgate aactagccag 540
caccatagtt ctccggaagtc agactatcat gatlaactgtg ctgactgtgc tctctattgt 600
actaaccatg ccgatgttta ggtgaaatta gctcacttg gcttcaacca tcttggtatc 660
cagttatcct cactgaattg agatttcttg ctccagtgtc agccattccc acataatttc 720
tgacctacag aggtgagggg tcataatagct ctccaaggat gctggtactc ccttcacaaa 780
ttcatttctc ctgttgtagt gaaaggtgcg cctctcggag cctccraggg tgggtgtgca 840
ggtcacatg atgaatgtat gatcgtgttc ccattaccca aaguccttaa atccctcatg 900
ctcagtaacc cagggcaggt ctagratctc ttcatttagt gtatgctgtc ccttcattgca 960
accacctcag gactcctgga ttctctgct agttgagctc ctgcatgtgt cctccttggg 1020
gaggtgaggg agaggggccc tggttcaatg ggalctgtgc agttgttaca ccttaggtgc 1080
tlaataaaca gaagctgtga tgttaaaaaa aaaaaaaa 1119

```

&lt;210&gt; 178

&lt;211&gt; 164

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; VARIANT

&lt;222&gt; (1) ... (164)

&lt;223&gt; Xaa - Any Amino Acid

&lt;400&gt; 178

```

Met. Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln Trp
1          5          10          15
Val Leu Ser Ala Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu
20        25        30
Gly Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val
35        40        45
Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro Leu Leu
50        55        60
Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu Ser
65        70        75        80
Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala Gly
85        90        95
Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Asp Ala Val
100       105       110
Ile Ala Ile Gln Ser Xaa Thr Val Gly Gly Trp Glu Cys Glu Lys Leu
115       120       125
Ser Gln Pro Trp Gln Gly Cys Thr Ile Ser Ala Thr Ser Ser Ala Arg
130       135       140
Thr Ser Cys Cys Ile Leu Thr Gly Cys Ser Leu Leu Leu Thr Ala Ser
145       150       155       160
Pro Gly Thr Leu

```

&lt;210&gt; 179

&lt;211&gt; 250

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 179

```

ctggagtgc tgggtatttc aagccctcgc aggaagcaga atgcaccttc tgaggcactc 60
ccagclgccc ccggccgggg gatgcggggc tgggagcacc ctggccgggc tgtgattgct 120
gcccggcact gttcattctca gttttctgt ccttttgctc ccggcaaggc cttctgctga 180
aagttcact utggagcctg atgtcttacc gaataaaggc cccatgctcc acccgaaaaa 240
aaaaaanaaa 250

```

<210> 180  
 <211> 202  
 <212> DNA  
 <213> Homo sapien

```

<400> 180
actagtcacg tgtggcgaaa ttccattgtg ttggggccaa cacaatggct acctttaacc 60
tcacccggac ccggcccttg ccgtgcgcc aogctgctgc taacgacagt atgatgctta 120
ctctgctact cggaaaactat ttttatgtaa ttatglatg cttctctgtc tataaatgcc 180
tgatttaaaa aaaaaaanaaa aa 202

```

<210> 181  
 <211> 558  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(558)  
 <223> n = A,T,C or G

```

<400> 181
cccytttktc naggtttkkg agacacccc agacctwaan ctgtgtcaca gaattcyngg 60
aatgtttagg cagtgcctagc aatttcytcg taatgattct gttattactt tccctnattct 120
ctattcctct ttctctctga gattaatgaa gttgaaatt gaggtggela aatacaaaa 180
ggtagtgtga tagtataagt atclaaagtgc agatgaaagt gtgttatata tatccattca 240
aaattatgca agttcgtaat tactcagggc taactaaatt accttaatal gctgttgaa 300
ctactctgtt ccttggtctag aaaaaattat aaacaggact ttgttagttt ggggaagccaa 360
attgataata ttctatgctt taazagtttg gctatacata aattattaag aaatatggaw 420
ttttattccc aggaatattg kgtttcattt atgaatatca cscrygatag awgtwtgagt 480
aaaaycagtt ttggtwaaata ygtwaatatg tcmteaataa acaakgcttl gaattatttc 540
caaaaaaaa aaaaaaaa 558

```

<210> 182  
 <211> 479  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(479)  
 <223> n = A,T,C or G

```

<400> 182
acagggwttk ggggatgcta agccccrga rwtgcttga tccaacctg gcttwtcttc 60
agaggggaaa atggggccta gaagktacag mscatytagy tggtyogmtg gcacccctgg 120
cttcacacag artcccgagt agctgggact acagacacac agtcaactga gcaggccctg 180
ttwgcaattc aogttgcac ctcccaetta aacattcttc atatgtgatg tctttagtca 240
ctaaggttaa acttccccc ccagaaaagg caacttagat aaaatcttag agtactttca 300

```

taetmttcta agtctctctc cagctctcat kkgagtcctm cytgggggtt gatagggaant.	360
ntctcttggc ttctctcaaa aartctctat ycatctcatg tthaalttgg tarcgacera	420
awtgstgaca aaattaaaat gtctctgctt marctttaaa aaaaaa aaaaaa	479

&lt;210&gt; 183

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 183

agggcgggagc agaaagctaaa gccaaagccr aagaagagty gcagtgccag caetgggtgcc	60
agtaccagta ccaataacag tgcagtgccr agtgccagca cragtgggtg cttcagtgct	120
gggtgccagcc tgacggccar tctcacattt gggtctcttc ctggcctctg tggagctggg	180
gccagracca gtggcagctc tgggtgctgt gggtctcttc acagtgaga ttttagatat	240
tgttaatcct gccagtgctt ctcttcagc cagggtgcat cctcagaaac ctactcaaca	300
cagcactcta ggcagccact atcaatcaat tgaagttagc actctgcaat aratctattt	360
gccatttcaa aaaaaa aaaa	384

&lt;210&gt; 184

&lt;211&gt; 496

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (496)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 184

accgaattgg gaccgctggc ttataagcga tcatgttynt ccrctatkar ctcaecgagc	60
agggagatcg agtctatarg ctgaagaaat ttgacccgat gggacaacag adctgctrag	120
cccatcctgc tgggtctctc ccagatgaca aatactcttg acccgaate accatcaaga	180
aacgcttcaa ggtgctcatg anccagcaac cggcgcctgt cctctgaggg tcccttaaac	240
tgatgtcttt tctgccacct gttacccctc ggagactcgg taaccgaact ctteggactg	300
tgagccctga tgcctttttg ccagccatac tctttggcat ccagtctctc gtygogattg	360
attatgcttg tgtgaggcaa tcatgggtgg atccccata aagggaacac atttgacttt	420
tttttctcat attttaatt actacmagaw tattwmagaw waaatgawt gaaaaactst	480
taaaaaaaa aaaaaa	496

&lt;210&gt; 185

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 185

gctggtagcc tatggcgkgs cccacggagg ggtcctgag gccacggcac agtgacttcc	60
caagtatcyt ggcsgcgtc ttctacrgtc cctacctgca gatcttcggg cagatcccc	120
aggaggacat ggacgtggcc ctcatygagc acagcaactg ytcghcggag cccggtctct	180
gggcacaccc tcttggggcc caggcgggca cctgctctc caagtatgcc aacttggctg	240
tggtgctgct cctngtcac tctctgctcg tggcgaacat cctgctggg aacttggctc	300
ttgcatgtt cagttacaca ttcggcaag tacagggcaa cagcagctct tcttgggag	360
gcgcagcgtt accgctcat cgg	384

&lt;210&gt; 186

&lt;211&gt; 577

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(577)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 186

gagttagctc	ctccacaacc	ttgatgaggt	cgctcgacgt	ggcctctcgc	ttcataccgc	60
tnccatcgct	ataatgtagg	tttggcccca	cytcctggca	tcttggggcg	gcntaatatt	120
ccaggaaact	ctcaatcaag	tcacccgtcg	tgaaccctgt	gggctgggtc	tgtcttcgcg	180
tcgggtgtaa	aggatctccc	agaaggagtg	ctcgatcttc	cccacacttt	tgatyaacttt	240
attgagtcga	ttctgcatgt	ccagcaggag	gttgtaccag	ctctctgaca	gtgaggtcac	300
cagcccatc	atgcgcttga	mcgtgccgaa	garcacagag	ccttgtgtgg	gggkkgaggt	360
ctcaccaga	ttctgcattc	ccagagagag	gtggcaaaag	acattgacaa	ctcgcgccag	420
gtggaaaaag	amcamctcct	ggargtgctn	gcgcctcttc	gtcmgttggg	ggcagcgctw	480
tcccttttgac	acacaaaaca	gttaaaaggca	tttccagccc	ccagaaantt	gtcatcatcc	540
aagatntcgc	acagcactna	tccagtggg	attaaat			577

&lt;210&gt; 187

&lt;211&gt; 534

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(534)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 187

aacatcttcc	tgtatcatgc	tgtgtaatat	cgatccgcatn	ttgtctggtg	agaatycatw	60
actkggaaag	gmaacattaa	agcctggaca	ctgggtattaa	aattcacaat	atgcacacact	120
ltaaacagtg	tgtcaatctg	ctccryynac	tttgcctaca	ccagctctggg	aakaagggtg	180
tgccttattc	acacctgtta	aaaggcgct	aagcattttt	gattcaacat	cttttttttt	240
gacacaaagtc	cgaaaaaagc	aaaagttaac	agttatyaat	ttgttagcca	attcaacttc	300
ttcatgggac	aggcccatyt	gatttaaaaa	gcaaatlgca	taatatggag	cttyggggagc	360
tgatatttga	gcggaagaqt	agcctttcta	cttcaccaga	cacaactccc	ttcatattg	420
ggatgttnac	naaagtwatg	tctctwacag	atgggatgct	tttgtggcaa	ttctgtcttg	480
aggatctccc	agtttattta	ccacttgca	aagaaggcgt	tttcttctc	aggr	534

&lt;210&gt; 188

&lt;211&gt; 761

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(761)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 188

agaaaccagt	atctctnaaa	acaaacctctc	ataccttgtg	gaacctaat	tgtgtgagtg	60
tgtgtgtgag	cgcatattat	atagacaggc	acatcttttt	tactttttgtg	aaagcttatg	120
cctctttggg	atctatatct	gtgaaagctt	taattgatctg	ccatnatgtc	ttggggagct	180



```

ttgtcttcty tgtaatggt actagagaaa acacctatnt tatgagtcuu tctagttngt. 240
tttattcgac atgaaggaaa ttccagatn acacactna caaactctcc ctkgackarg 300
ggggacccag auaagccaaa ctgamcataa raaacatwa cctgggtgag arttgcataa 360
acagaatwr ggtagtatat tgaatnacg catcattaaa rmggttwktt wttctccctt 420
gcacaaaaeca tgtacngact tccggttgag taatgcccag ttgtcttctt catnataaaa 480
cttgcccttc attacatggt tnaaagtgt gtggtgggnc aaaaatattga aatgatggaa 540
ctgactgala aagutgtaca aataagcagt gtgcctaaac agcaacacag taatgttgac 600
atgcttaatt cacaaatgct aatltcatta taaatgtttg ctcaaatata ctctgaacta 660
ttttctctgt ttccagagc tgagatntta gcttttatgt agtatnaagt gaaaaantac 720
gaaantaata acattgaaga aaaaaaanaa aaaaaaaa a 761

```

&lt;210&gt; 189

&lt;211&gt; 482

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(482)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 189

```

tttttttttt ttgcccgatn ctactatttt attgcaggan gtgggggtgt atgcaaccga 60
caccgggggt atnagaagca agaagggaagg agggaggggca cagcccttg ctgagcaaca 120
aagccgcttg ctgcttctc tgtctgtctc ctggtgtagg ccatgggga gaccttcccc 180
aaggcagggg ccaccagtcu aggggtggga atcagggggg tgggagtggt acataagaag 240
tgatagggau aggcaccccg gtacagacuu ctgggtctct gacaggtnga tttegaccag 300
gtcattgtgc cctgcccagc caccgcgtan atctggaaaa gacagaatgc ttctcttttc 360
aatltgggt ngtcutngaa ngggcatttt tcaaatntng gctnggtctt ggtacncttg 420
gttcggccca gctcncgtc caaaantcat tcacccnctt ccaaatgtgt tgcngnncr 480
cc

```

&lt;210&gt; 190

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(471)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 190

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tttttttttt ttttbaaca gtttttraca acaaaattta ttagaagaat agtgggtttg 60
aaaactcttg cctccagtga gaactaccat acaccacatt acagctngga atgtncctca 120
aatgtctggt caaatgatgc aatggaaaca ttcaatctta cacatgcacg aaagaacaag 180
cgctttttgac atacaatgca caaaaaaaa aggggggggg gacacatgg attaaaattt 240
taagtactca taccatacat taagacacag ttctagtcca gtcaaaaate agaactgcnt 300
tgaaaaattt catgtatgca atcaaaccaa agaacttntt tggtagtcat gantnctcta 360
ctacatunac cttgatcatt gccaggaacn aaaggttnaa oncacnngt acaaaaaana 420
tctgtaattn anttcaact cgtacngaa aeatntntnt tataacttcc c 471

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&lt;210&gt; 191

&lt;211&gt; 402

&lt;212&gt; DNA

<213> Homo sapien

<220>

<221> Misc\_feature

<222> (1) ... (402)

<223> n = A,T,C or G

<400> 191

gagggattga	aggctctgtt	tastgtoggn	ctgttcagcc	arcaaactcta	acaagttgct	60
gtcttccact	cactgtctgt	aagcttttta	acccagacwg	tatcttcata	aatagaacaa	120
attcttccac	agtcacatct	tctaggacct	ttttggatto	agttagtata	agctcttcca	180
cttcccttgt	taagacttca	tctggtaaa	tcttaagttt	tgtagaaagg	aattyaattg	240
ctcgttctct	aacaatgtcc	tctccttgaa	gtatttggct	gaacaaccca	cctaaagtcc	300
ctttgtgcat	ccatttttaa	tatacttaet	agggcattgk	tncaactaggt	taaattctgc	360
aagagtcato	tgtctgcaaa	agttgogtta	gtatctctgc	ca		402

<210> 192

<211> 601

<212> DNA

<213> Homo sapien

<220>

<221> Misc\_feature

<222> (1) ... (601)

<223> n = A,T,C or G

<400> 192

gagctcgggt	ccataatct	ttgtctgagg	gcagcacaca	tatncagtgc	catggnaact	60
ggtctacccc	acatgggagg	agcatgcrct	agntatataa	ggtcatttcc	tgagccagac	120
atgcytyttt	gaytaccgtg	tgccaaagtcc	tggtgattcl	yaacacacyt	ccatcccgyt	180
cttttgtgga	aaaactggca	cttkctctga	actaggarga	catcacttcc	aaattcaccc	240
acgagacact	tgaagggtgt	aacaaagcga	ytcttgcaat	gctttttgtc	cctccggcac	300
caattgtcaa	tactaacccg	ctggttttgc	tccatcacat	ttgtgatctg	tgtctctgga	360
tacatctcct	gcagtaactg	aagaacttct	tcttttgttt	caaaagcacc	tcttggtgcc	420
tgtcagatca	ggttcccatc	tccagctcgg	aatgttcaca	tggcatattt	waattcccaa	480
aaaacattgc	gattttaggg	tcagcaacag	caaatcctgt	tccggcattg	gctgcaagag	540
cctcagatga	gcccggccag	gccaaaggcg	gcccctgag	ccccaccagg	agcagaagca	600
s						601

<210> 193

<211> 608

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1) ... (608)

<223> n = A,T,C or G

<400> 193

atcacagcca	natccaccca	cgaagatgag	cttgtttgact	gagaacctga	tgaggctcact	60
ggtcccgtcg	tagcccccag	gactctccac	ctgctggaag	cggttgatgc	tgcactcytt	120
cccaacgcag	gcagmagcgg	gscgggtcaa	tgaactccay	tctgsgcttg	gggtkgacgg	180
tkaagtgcag	gaagaggtcg	accacctngc	ggtccacacg	gatccccgac	tgtgcgggac	240
ctgcagcga	actcctcgat	ggtcattgag	gggaagcga	tgaggccacg	ggccttgccc	300

agaaccttcc	gcctgttctc	tggugtcacc	tgcagctgct	gcegetgaca	ctcgguctcg	360
gaccagcgga	caaacaggct	tgaacagccg	cacctcacgg	atgcccagtg	tgtcgrgctc	420
caggammjgc	accragctgt	ccaggtcaat	gtcgggtgaag	ccctccggcg	gtralgycgt	480
ctgcagctgt	tltgtcgatg	ttctccaggg	acaggtctgc	cagctgaggc	tcatcgaaga	540
gtcgcgcttg	cgtgagcagc	atgaaggcgt	tgtcggtctg	cagttcttct	tcagggaactc	600
cacgcaac						608

&lt;210&gt; 194

&lt;211&gt; 392

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(392)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 194

gaacggctgg	accttgcttc	gcatttgtgt	tgtctggcagg	gaataccttg	gcaaggagyt	60
ccagtcagag	cagccccaga	ccgtgcgcgc	cngaggttaa	gcctgcctct	ggccttcccc	120
tccgctcaa	tgcagaacca	gtagtgggag	cactgtgttc	agagttcaga	gtgaacactg	180
tttgatttta	cttgggaatt	tccctctgta	tatagctttt	cccaatgcta	atttccaaac	240
aacaacaca	aaataacatg	tttgctgttc	aagttgtata	aaagtaggtg	attctgtatt	300
ttaagaaat	attactgtta	catatactgc	ttgcaatttc	tytattttatt	gkctctatgg	360
aaataaatat	agttatttaa	ggttgttaat	cc			392

&lt;210&gt; 195

&lt;211&gt; 502

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(502)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 195

ccattkgagg	ggtkaggkyc	cagttycgcg	gtggaagaaa	caggccaggga	gaagtgcgtg	60
ccagctgag	gcagatgttc	ccacagtga	ccccagagcc	stgggatata	gtytctgacc	120
cctcncaagg	aaagaccaca	ttctgggggc	atgggtctga	ggcraggacc	tagaggcacc	180
aagggaaagg	cccattccgg	ggatgttccc	cgaggaggga	gggaaggggc	tctgtgtgcc	240
cccaaggagg	aaggggcccc	gagtccctgg	atcagacacc	ccttcacgtg	tatccccaca	300
caantgcaag	ctcacccaag	tccccctcca	gtcccccttc	ataccacctg	amcyggccact	360
gscscacacc	cacccagagc	acgccacccg	ccatggggar	tgtgctcaag	gartcgcnng	420
gcacgctgga	catctngtcc	cagaaggggg	cagaactctcc	aatagangga	ctgarcattt	480
gctnanaaaa	aaaaananaa	aa				502

&lt;210&gt; 196

&lt;211&gt; 665

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(665)

<223> n = A,T,C or G

<400> 196

ggttacttgg	tctcattgcc	accacttagt	ggatgbcatt	tagaaccatt	ttgtctgccc	60
ccctcgaag	ccttgcgcag	agcggacttt	gtatttgttg	gaggatcact	gctgaatbtt	120
wagctgtttk	gagttgaltt	gcaccactgc	acccacacat	tcaatatgaa	aacyawttga	180
actwatttat	tatcttgtga	aaagtataac	aatgaaaatt	ttgttcatac	tgtatrkac	240
aagtatgatg	aaaagcaawa	gatatatatt	cttttattat	gttaaattac	gattgccatt	300
actaatcggc	aaatgttggg	gtgfatgttc	ttttcacagt	aatatattgc	ttttgttaact	360
tracttgggt	atttttattg	aaatgattta	caaaattott	aatttaagar	aatggatatg	420
watatttatt	tcatttaatt	ctttcctkgt	ttacgtwaat	tttgaaaaga	wtgnatgatt	480
ttttgacaga	aatcgatcct	gatgctgttg	aagtatgttg	acccacatcc	ctatgggttt	540
ttcttagant	gtataaaggt	tgtagcccat	unaacttcaa	agaaaataat	gacacatac	600
tttgcaatra	ggctgaaatg	tggcatgctn	ttctaatcc	aatcttataa	actagcaaan	660
aagtg						665

<210> 197

<211> 492

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1) ... (492)

<223> n = A,T,C or G

<400> 197

ttttttttct	ttttttttgc	aggaaggatt	ccattttattg	tggatgcatt	ttcacatat	60
atgttttattg	gagcgatcca	ttatcagtga	aaagtatcaa	gtgtttataa	natttttagg	120
aaggcagatt	cccgaaacat	gctngtongc	ttgcagtttt	acctcgtaaa	gatnacagag	180
aattatagtc	naaccagtaa	acnagggaatt	tacttttcaa	agatttaaat	ccaaactgaa	240
caaaattcta	ccctgaact	tactccatcc	aatatttggg	ataanagtcc	gcagtgtatc	300
atctctttct	gaacttttag	ttttctagaa	anatatgtaa	tgtgtatcag	gaagagctct	360
tgttcaaaag	tacaacnaag	caatgtttcc	ttaccatagg	ccttaattcc	aattttgatc	420
catttcaatc	ccatcacggg	agtcactgct	acctgggaca	cttgtatttt	gttcatnctg	480
ancttggtt	aa					492

<210> 198

<211> 478

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1) ... (478)

<223> n = A,T,C or G

<400> 198

ttttttttg	atttcantct	gtannaanta	ttttcattat	gtttattana	aaatatnaa	60
tgtntccarn	acaaatcatn	ttacntnagt	aaggaggccan	ctacattgtc	caacatarac	120
tgagtatatt	ttgaaaagga	caagttttaa	gtanacnaat	attgcoganc	atancacatt	180
tatacatggc	ttgattgata	tttagcacag	canaaactga	gtgagttacc	agaaanaaat	240
netatatgtc	aatengattt	aagatacaaa	acagatctta	tggtacatan	catntgtag	300
gagttgtggc	tttatgttta	ctgaaagtra	atgcagttcc	tgtacaaaga	gatggccgta	360
agcattctag	tacctetact	ccatgggtta	gattcgtaaa	cttatgttta	catatgttca	420

gggtcagat tctgtttagt naanttatgg agaggtccan gagaaaaatt tgatncaa 478

<210> 199

<211> 482

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> {1}...{482}

<223> n = A,T,C or G

<400> 199

agtgaattgt	cttccncaa	aacctctga	tcaagtttgt	ggcaactgaca	atcagaccta	60
tgctagttcc	tgtcatctac	togctartaa	atgcagactg	gaggggacca	aaaaggggca	120
tcaactccag	ctggattatt	tggagacctg	caaatctatt	cttacttgta	cggactttga	180
agtgattcag	tttctctac	gaatgagaga	ctggctcag	aatactctca	tgcagcttta	240
tgaagccnac	tctgaacacg	ctggttatct	agatgagaa	ncagagaaat	aaagtcnaga	300
aaatttacct	ggangaaaag	aggcttngg	ctggggacca	tccattgaa	cttctctta	360
anggacttta	agaaanaact	accaratgt	tgtngtatcc	tggtagcngg	cgtttantg	420
aactngacn	ncaccttnt	ggatatanant	cttgacngn	tctgaactt	gtctctctgc	480
ga						482

<210> 200

<211> 270

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> {1}...{270}

<223> n = A,T,C or G

<400> 200

gggcggcaag	tgcaactcca	gctggggcgg	tgcggagcaa	gattctgccn	gcagttggtc	60
cgactgcgac	gacggcgggc	ggcagctcg	caggtgcagc	gcggggccct	gggtctctgc	120
agggtgagc	tgacgcccga	gaggtcgtgt	cagctccccc	gaccttgacy	cgtcggggga	180
cagcgggac	agagcccggt	gaagcggga	ggcctcgggg	agccctcggg	gaagggcggc	240
ccgagagata	cgcaggtgcn	ggtggccgcc				270

<210> 201

<211> 419

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> {1}...{419}

<223> n = A,T,C or G

<400> 201

ttttttttt	ttttggaac	tactgcgagc	acagcaggtc	agcaacaagt	tlattttgca	60
gtatgcaagg	taacagggta	gggcatgggt	acatgttcag	gtcaacttcc	tttgttgtgg	120
ttgattgggt	tgtctttatg	ggggcggggt	ggggtagggg	aaancgaagc	anaantaaca	180
tggagtgggt	gcacctccc	tttagaacct	ggttacnaaa	gcttggggca	gttcaactgg	240

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tctgtgaccg tcatttttctt gacatcaabg ttattagaag tcaggatato ttttagagag 300
tccactgtnt ctggaggag attagggttt ctggccaaa tccancaaa atccacntga 360
aaaagttagg tcatncangt acngaatccc gaggccatan ttctcatant cgggtggcaa 419

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&lt;210&gt; 202

&lt;211&gt; 509

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(509)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 202

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tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt 60
tggcactttaa tccatttttta ttccaaaatg tctacaaant ttnaathncuc cattatacng 120
gttttttttc aaaaatctaaa ntttattcaa atotnagcca aatttccttac ncaaatmaa 180
tarnncnaaa aatcaaaaat ataentntct ttccagcaaac ttngtccat aatttataaa 240
aatatatacg gctgggtgtt tcaaaagtaca attatcttaa caatgcaaac atttttttaa 300
ggaactaaaa caaaaaaaa cactnccgca aaggttaag ggaacaaaca attcctttta 360
caacancnc nattataaaa atcabatctc aatctttag ggaatatata ctcccaacng 420
ggatcttaac ttttactnca ctgtgttat tttttttaa ccattgtntt gggcccaaca 480
caatggnaat nccnccncc ttgactagt 509

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&lt;210&gt; 203

&lt;211&gt; 583

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(583)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 203

```

tttttttttt ttttttttga cccccctctt ataaaaaaca agtkaccatt ttattttact 60
tacacatatt ttttttttaa ttggtattag atattcaaaa ggcagctttt aaaaatcaaac 120
taaatggaaa ctgccttaga tacataattc ttaggaaata gcttataaac tgcctaaagt 180
gaaaatcttc tctagctctt ttgactgtaa attttctgact ctgtataaac atccaaatcc 240
attttctctg tctttaaaat tatctaactt tcccattttt tccctatttc aagtcatttt 300
gottctctag cctcatttcc tagctcttat ctactattag taagtggctt ttttccataa 360
agggaaaaaa ggaagagana atggcaacaa aaacaaacat ttctatttca ttttctacc 420
tacgttaata aatagcatt ttgtgaagcc agtcataaag aaggcttaga tctttttatg 480
tccatttttag tcaataaacc atatcnaaag tgcacgaatg caaaagggtt gtgaacattt 540
attcaaaagc taatataaga tatttcacat aatcatcttt ctg 583

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&lt;210&gt; 204

&lt;211&gt; 589

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(589)

<223> n = A,T,U or G

<400> 204

tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	60
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	120
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	180
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	240
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	300
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	360
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	420
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	480
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	540
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	589

<210> 205

<211> 545

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(545)

<223> n = A,T,C or G

<400> 205

tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	60
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	120
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	180
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	240
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	300
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	360
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tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	480
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	540
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	545

<210> 206

<211> 487

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(487)

<223> n = A,T,C or G

<400> 206

tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	60
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	120
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	180
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	240
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	300
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	360
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	420
tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	480

ttcamaa

487

&lt;210&gt; 207

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (332)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 207

tgaattggct aaaaagactg atttttanaa clagcaactc ttatttcttc cctttaaaaa	60
lacataquut taaatcccaa atccttctta aagacctgac agcttgagaa ggtcactact	120
gcatttatag gaccttctg tggttctgct gttacttttg aantctgaca atccttgana	180
atcctttgcat gcagggaggg taagaaggtat tggattttca cagaggaana acacagcgca	240
gaaatgaagg ggccagggtt actgagcttg tccactggag ggctcatggg tgggacatgg	300
aaaagaaggc agcctaggcc ctggggaguc ca	332

&lt;210&gt; 208

&lt;211&gt; 524

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (524)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 208

agggcgtggt gcagaggcgg ttactgtttt gtctcagtaa caataaatac aaaaagactg	60
gttgtgttcc ggcacctcc aaccacgaag ttgatttctc ttgtgtgcag agtgactgat	120
tttaaggac atggagcttg tcacaatgtc acaatgtcac agtgtgaagg gcacactcac	180
tcccggtga ttacattta gcaacacaca atagctcatg agtccatart tgaataact	240
tttggcagaa tecttnttga aacttgacga tgataactaa gatccagat atttcccaa	300
gtaaatagaa gtgggtcata atattaatta cctgttcaca tgggttcca tttaaaagtc	360
atgagcccg acactgacat caaactaagg caacttagac tcttcaccac cagtctgtcc	420
tgatcagca caggaggctg tcaccttgac caaattctca ccagtcact atctatcna	480
aaaccattac ctgattcact tgggtaatg caccacttg gtga	524

&lt;210&gt; 209

&lt;211&gt; 159

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 209

gggtgaggaa atccagagtt gccatggaga aaattccagt gtcagcattc ttgtctcttg	60
tggcctctc ctacactctg gccagagata ccacagtcac acctggagcc aaaaaggaca	120
caaaggactc tgaacccaaa ctgcccaga cctcttca	159

&lt;210&gt; 210

&lt;211&gt; 256

&lt;212&gt; DNA

&lt;213&gt; Homo sapien



<220>  
 <221> misc feature  
 <222> (1)...(256)  
 <223> n = A,T,C or G

<400> 210  
 actccctggc agacaaaggu agaggagaga gctctgttctg ttctgtgttg ttgaactgcc 60  
 actgaatttc ttcccaattg gactattaca tggcatttga gggactaatg gaaaaacata 120  
 tggggagatt ttanccaatt tangtntgtt aatgggagga ctggggcagg cggggagagat 180  
 ttgaggggtg naaatgggan ggctgggttg tcanatgaac agggacatag gaggtaggca 240  
 ccaggatgct aaatca 256

<210> 211  
 <211> 264  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc feature  
 <222> (1)...(264)  
 <223> n = A,T,C or G

<400> 211  
 acattgtttt tttagataa agcattgaga gagctctctt taacttgaca caatggaagg 60  
 actggaacac ataccacat cttagtcttg agggataatt ttctgataaa gtcttgctgt 120  
 atattcaagc acatstgta tatattttt agtccatgt ttatagccta gtttaggaga 180  
 gggggagatac attcngaaag aggaactgaa gaaatactca agtnggaaaa cagaaaaaga 240  
 aaaaaggag caaatgaga gcct 264

<210> 212  
 <211> 328  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc feature  
 <222> (1)...(328)  
 <223> n = A,T,C or G

<400> 212  
 acccaaaaat ccaatgttga atatttggct tcattatttc canattcttt gattgtcaaa 60  
 ggatttaatg ttgtctcagc ttggycaatt cagttaggac ctagggatgc cggctgggag 120  
 gtttatatat gcagcaacaa tattcaagcg agcaacragg ttattgaatt tgcctggcag 180  
 tttaatttca ttccatttga ctggggatcc ttatcatcag ccagagagat tgaaatttta 240  
 cccctacnac tctttactct ctgganaggg ccagtgggtg tagctataag cttaggcaca 300  
 ttttttttct ctttatttct ttgtcaga 328

<210> 213  
 <211> 250  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc feature

&lt;222&gt; {1}...(250)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 213

acttatgagc agagcgacat atccnagtgt agactgaata aaactgaatt ctctccagtt	60
taaagcattg ctcaactgag ggatagaagt gaatgccagg agggaaagta agccaaaggt	120
cattatgcca aagganatat acatttcaat tctccaaact tcttctcat tccaagagtt	180
ttcaatattt gcataaact gctgataene catgttaana sacaaatate tctctnacct	240
tctcatcggt	250

&lt;210&gt; 214

&lt;211&gt; 444

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...(444)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 214

accagante caatgctgaa tatttggtct cattattccc agatcttttg attgtcaaag	60
gatttaattgt tctctcagct tgggcacttc agttaggacc taaggatgac agccggagg	120
tttatatatg cagcaacaa. attcaagcgc gacaacaggt tattgaactt gcccgccagt	180
tgaatttcak tcccuttgac ttgggatcct tatcatcagc canagagatt gaaaatttac	240
ccctacgact ctttactctc tggagagggc cagtgtgtgt agctataagc ttggccacat	300
tttttttct tttattcctt tgtcagagat gogattcacc calatgotan aaaccacag	360
agtgcctttt acaaaattcc tataganatt gtgaatcaaa ccttacctat agttgccatt	420
actttgctct cctaatata cctc	444

&lt;210&gt; 215

&lt;211&gt; 366

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...(366)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 215

acttatgagc agagcgacat atccaaagtgt aaactgaata aaactgaatt ctctccagtt	60
taaagcattg ctcaactgag ggatagaagt gaatgccagg agggaaagta agccaaaggt	120
cattatgcca aagganatat acatttcaat tctccaaact tcttctcat tccaagagtt	180
ttcaatattt gcataaact gctgataaag catgttgaga sacaaatate tctctgaact	240
tctcatcggt aagcagagggc tctaggcaac atggaccata gcgaanaaaa aacttagtaa	300
tccaagctgt tttctacact gtaaccaggt ttccaaccaa ggtgggaate tcttataact	360
ggtgcc	366

&lt;210&gt; 216

&lt;211&gt; 260

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

<221> misc\_feature  
 <222> (1)...(260)  
 <223> n = A,T,C or G

<400> 216

ctgtataaac	agaaatccac	tgcangaggg	agggccgggc	caggagaatc	tccgcttgtc	60
caayacaggg	gcctaaggag	ggctccacaa	ctgctnntaa	gggtcttttc	attcttttat	120
taataaaaag	tnnaaaaggc	ctctctctca	cttttttccc	ttnggctgga	aaatttaaaa	180
atcaaatatt	tccnaagtt	ntcaagctat	catatatact	ntatcctgaa	aaagcaaat	240
aattcttcc	tccctcttt					260

<210> 217  
 <211> 262  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(262)  
 <223> n = A,T,C or G

<400> 217

acctaagtg	gtangttan	aaatgttata	atttcaggaa	naggaacgca	tataattgta	60
tcttgctat	aattttctat	tttaataagg	aaatagcaaa	ttggggtgga	gggaatgtag	120
ggcatctac	agtttgagca	aaatgcaatt	aaatgtggaa	ggacagcact	gaaaaatttt	180
atgaataatc	tgtatgatta	tctgtctcta	gagtagattt	ataattagcc	acttacccta	240
atctcttca	tgtctgtaa	gt				262

<210> 218  
 <211> 205  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1)...(205)  
 <223> n = A,T,C or G

<400> 218

accaaggtg	tgcattaccg	gaantggatc	aangacarca	tngtggccaa	ccctgagca	60
ccctatcaa	ctcccttttg	tgttaactt	ggaaccttgg	aatgacrag	gccaagactc	120
aggcctccc	agttctactg	acctttgtcc	ttangtntna	ngtccaggg	tgtaggaaa	180
anatacagc	agacacaggt	gtana				205

<210> 219  
 <211> 114  
 <212> DNA  
 <213> Homo sapien

<400> 219

tactgttttg	tctcagtaac	aataaataca	aaaagartgg	ttgtgttccg	gcccacatcc	60
accangaagt	tgattttct	tgtgtgcaga	gtgactgatt	taaaaggaca	tggg	114

<210> 220  
 <211> 93

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 220

actagccagc acaaaaggca gggtagcctg aattgcttcc tgcctcttbaa atttctctta 60  
 aaataagcat ttagtgcra gtcctactg agt 93

&lt;210&gt; 221

&lt;211&gt; 167

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{167}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 221

actangtgcg ggtgcgcaca aatatttgcg gatattccct tcactcttga ttccatgagg 60  
 tcttttgcgc agcctgtggc tctactgtag taagtctctg ctgatgagg gccagatgc 120  
 ccccaactac ctccctgac gctcccaaa aatcaccaa cctctgt 167

&lt;210&gt; 222

&lt;211&gt; 351

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 222

agggcgtggt ggggagggcg gtactgacct cattagtagg aggatgcatt ctggcaccoc 60  
 gttcttcacc tgcaccccaa tccttaaaag gccatcttgc ataaagtcaa caacagataa 120  
 atgtttgctg aattaaagga tggatgaaa aattaataa tgaatttttg cataatccaa 180  
 tttctctctt tatatttcta gaagaagttt ctttgagcct attagatccc ggggaatctt 240  
 taggtgagca tgaattagaga gcttgtaggl tgcctttaca tatctctggc atatttgagt 300  
 ctogtatcaa aacaatagat tggtaagggt ggtattattg cattgataag t 351

&lt;210&gt; 223

&lt;211&gt; 383

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...{383}

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 223

aaacacaca aacaaaaaa acattcttc attcagaaa attatcttag ggaactgatat 60  
 tggtaattat ggtcaattta atwrtttkt ggggcatttc cttaacattgt ctgacacga 120  
 ttaaaatgtc tgggcacaaa ttttgattt tatttgaga ctctctatca aagtaatgc 180  
 tgcacaagga agtctaagga attagttagt tttccmccac ttgtttggag tgtgctatc 240  
 taaaagattt tgaatttctg gaatgacact tatattttaa ctttggtagg ggaanagtt 300  
 ataggaccac agtcttcaat tctgatactt gtaaatteat ctttatttgc atttatttg 360  
 accattaagc tatatgttta aaa 383

&lt;210&gt; 224

<211> 320  
 <212> DNA  
 <213> Homo sapien

<400> 224

cccttgagg	cttcttgta	gaaatagta	cagttacac	caataggaa	aacaaaaa	60
aaaagtgtg	gacattgtg	tagggagtgt	gtacccctta	ctccccctc	aaaaaaat	120
ggatavctg	ttaaaggata	raagggcaat	atcttatcat	atgtttctaa	agaggaaggaa	180
gagaaaaaac	tactttctc	aatgggaagc	ccctaaagggt	gctttgatac	tgaaggacac	240
aaatgtggcc	gtccatctc	ctttaragtt	gcattgactt	gacacggtaa	ctgttgaggt	300
tttaractcm	gcattgtgac					320

<210> 225  
 <211> 1214  
 <212> DNA  
 <213> Homo sapien

<400> 225

gaggactgca	gccccactc	gaagccctgg	caggccggac	tggtcatgga	aaargacttg	60
ttctgctcgg	gggtcctgg	gcacccggc	tggtgtctgt	cagccgcaca	ctgtttccag	120
actcctaca	ccatcgggt	gggtctgac	agtccttgag	ccgaccaaga	gcccgggggc	180
cagatgggtg	aggtccagct	ctcgtacgg	cauccagagt	acaaacagac	cttgcctcgt	240
acagacctc	tgtccatcaa	gttggacgaa	tcctgtctcg	agtctgacac	ctcgcggagc	300
atcagcttg	cttcgagtg	ccctaccgg	gggaactctt	gctcgtttc	tggctggggc	360
ctgctggcga	acggcagaat	gcctaccgtg	ctgcagtggc	tgaacgtgtc	gggtggtgtc	420
gaggaggtct	gcagttaagc	ctatgacctg	ctgtaccacc	ccagcatgtt	ctgcgcgggc	480
ggagggcaag	accagaagg	ctcctgcaac	ggtgactctg	gggggcccc	gactctgcac	540
gggtacttgc	agggccttgc	gtttttcgga	aaagccccgt	gtggccaagt	tygcgtgcca	600
gggtgtctac	ccaaacctctg	caaattcccl	gagtggtatg	agaaaaacgt	ccaggccagt	660
taactctggg	gactggggaac	ccatgaatt	gaccccaaa	tacatctctg	gggaaggatt	720
caggaataat	tgtttccagc	ccctcctccc	tcaaggccag	gagtcagggc	ccccagcccc	780
tcctccctca	aaccaagggt	acagatcccc	agccccctct	ccctcagacc	caggagtcca	840
gacccccag	ccctctctcc	ctcagaccca	ggagtccagc	ccctcctccc	tcagacccag	900
gagtcagagc	ccccagccc	ctcctccctc	agacccaggg	gtccaggccc	ccaaacctct	960
ctccctcaga	ctcagaggtc	caagccccca	acccctcctt	ccccagaccc	agaggtccag	1020
gtccagagcc	ctcctcctc	agacccagcg	gtccaatgac	acctagactc	tccctgtaca	1080
cagtgccccc	ttgtggcag	tgacccaac	cttaccagtt	ggttttctat	tttttgtccc	1140
tttcccttag	atcagaaat	aaagtctaat	agaagcgcac	aaaaaasaaa	aaaaaasaaa	1200
aaaaaaaaaa	aaaa					1214

<210> 226  
 <211> 119  
 <212> DNA  
 <213> Homo sapien

<400> 226

acccagtatg	tgcaggagga	cggaacccca	tgtgacagcc	cactccacca	gggttcccaa	60
agaacctggc	ccagtctaaa	tcattcatcc	tgaagtgagg	aaataacag	ataaccagt	119

<210> 227  
 <211> 818  
 <212> DNA  
 <213> Homo sapien

<400> 227

acaattcata	gggacgacca	atgaggacag	ggatgaacc	eggctctccc	ccagccctga	60
tttttctac	atatgggggc	ccttttcatt	ctttgcaaaa	acactgggtt	ttctgagaac	120
acggacgggt	cttagcaca	tttgtgaat	ctgtgtaraa	ccgggctttg	caggggaggt	180
aatttctct	ctctggagga	aaggtggtga	ttgaragga	gggagacagt	gacaaaggct	240
gagaagggc	cgctcggcct	tctctgaacc	agggtggaac	ggcagacccc	tgaaaaagaa	300
agttgtcccc	ttccaatcag	ccaattctga	gaaccccct	ctaaattctt	actggaaaag	360
agggcctcct	caggagcagt	ccaagagbtt	tcnaagataa	cgtgacaaat	accatctaga	420
ggaaagggtg	cacctctagg	ayagaagccg	agagctttaa	tctggctcgt	ttccagagaca	480
acctgctggc	tgtcttggga	tgcgcccagc	ctttgagagg	ccactacccc	atgaacttct	540
gcctatccat	ggacatgaag	ctgaggacac	tgggcttcaa	cactgagttg	tcattgagagg	600
gacaggtctt	gcctcgaagc	cggtctgagg	uagcaaccac	tctcctcccc	tttctcagcc	660
aaagccattc	ccacaaatcc	agacatatac	atgaagcaac	gagaccctaa	cagtttggct	720
caagaggata	tgaggactgt	ctcagccctg	ctttgggctg	acaccatgca	cacacacaa	780
gtccacttct	aggttttcag	cctagatggg	agtctgtg			818

<210> 228

<211> 744

<212> DNA

<213> Homo sapien

<400> 228

actggagaca	ctgttgaact	tgatcaagac	ccagaccacc	ccaggtctcc	ttctgtgggt	60
gtcatgacgt	ttgacatacc	tttggaaaga	gcctcctcct	tggaagatgg	aagaccgtgt	120
tcgtggccga	cctggcctct	cctggcctgt	ttcttaagat	gggaggtcac	atttcaatgg	180
taggaaaagt	ggcttcgtac	aatagaagag	cagtcactgt	ggaactacca	aatggcgaga	240
tgtctgggtg	acattggggg	gctttggggt	aaaagattta	tgagcgaact	attctctggc	300
accagattct	aggccagttt	gttccactga	agcttttccc	acagcagtrc	acctctgcag	360
gctggcagct	gaatgggttg	ccggtgggtc	tatggcaaga	tcacactgag	atcgatgggt	420
gagaagggta	ggatgcttgt	ctagtgttct	tagctgtcac	gttgggtcct	tcaggtttgg	480
ccagacgggt	ttggccactc	cctttcaaaa	ccagggcgcc	ctcctgggtg	cagtgcaccg	540
ccgtgggtat	ccttggccca	ttccagcagt	cccagttatg	catttcaagt	ttggggtttg	600
ttcttttctg	taatgttctt	ctgtgttgtc	agctgtcttc	atttccctgg	ctaaagcagc	660
ttgggagata	tgaccagag	atccactcct	taagaaccag	tggcyaaga	cactttcttt	720
cttcaactct	aagttagctgg	tgg				744

<210> 229

<211> 300

<212> DNA

<213> Homo sapien

<400> 229

cgagttctgg	ttttgtctat	aaagtttgat	ccctcctttt	ctcatccaaa	tcattgtgaac	60
cattacacat	cgaaataaaa	gaaaggtggc	agacttgccc	acggccaggc	tgacatgtgc	120
tgcaggggtg	ttgtttttta	attattatgt	ttagaagcgt	cacccacagt	ccctgttaat	180
ttgtatgtga	cagccaactc	tgayaaaggt	ctatttttcc	acctgcagag	gatccagtrt	240
cactaggctc	ctccttgccc	tcacactgga	gtctccgcca	gtgtgggtgc	ccactgaact	300

<210> 230

<211> 301

<212> DNA

<213> Homo sapien

<400> 230

cagcagaaca	aatacaaat	tgagaggtgc	aaagatctca	taaaatctat	actgagggaat	60
gagcgacagt	tcnaaggagga	gaagcttcca	gagcagctca	agcaagctga	ggagctcagg	120

caatataaag tccctgggtca cactcaggga cagagagctga cccagtttaag ggaagaagttg 180  
 cgggaaggga gagatgcctc cctctcattg aatgagcatc tccaggccct cctcactccg 240  
 gatgaacgg acaagtcaca ggggcaggac ctcccaagaa cagacctcgg ccguyaccac 300  
 g 301

<210> 231  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 231  
 gcaagcacgc tggcaaatct ctgtcaggtc agctccagag aagccattag tcatrtttag 60  
 cagggaactcc aagtcacat ccttggcaac tggggacttg cgcaggttag ccttgaggat 120  
 ggcaacacgg gacttctcat caggaagtgg gatgtagatg agctgatcaa gacggccagg 180  
 tctgaggatg guaggatcaa tgatgtcagg ccggttggta ccgccaatga tgaacacatt 240  
 tttttttgtg gacatgccat ccatttctgt caggatctgg ttgatgactc ggtrcagcagc 300  
 c 301

<210> 232  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 232  
 agtaggtatt tcgtgagaag ttcaaacacca aaactgggac atagttctcc ttcaagtgtt 60  
 ggcgaacagcg gggcttccctg attctggaat ataactttgt gtaaattaac agccacctat 120  
 agaagagtc atctgctgtg aaggaagagac agagaactct gggctccgtc gtccctgtcc 180  
 cgtgctgtac caagtgtctg tgcagccctg ttacctgttc ccaactgaaa tctggctaat 240  
 gctctgtgt atcattctg attctgacaa tcaatcaatc atgggctag agcactgact 300  
 g 301

<210> 233  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 233  
 atgactgact tccagtaag gctctctaa gggtaagtag gaggatcac aggatttgag 60  
 atgctaagga cccagagatc gtttgatcca accctcttat ttccagaggg gaaatgggg 120  
 cctagaagtc acagagcatc tagctggtgc gctggcacc ctggcctcac acagactccc 180  
 gagttagctg gactacagga acacagtcac tgaagcagga cctgttagca attctatgcg 240  
 tacaattaa cctgagatga gtagagactt tattgagaaa gcaagagaaa atcctatcaa 300  
 c 301

<210> 234  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 234  
 aggtcctaca catcgagact catccatgat tgaatgaat ttaaaatla caagcaanga 60  
 catthtattc atcatgatgc ttctttttgt ttctttttt cgttttttt ttttttttt 120  
 tcaatttcag caacatactt ctcaatttct tcaaggattta aaatctttag ggattgatct 180  
 cgcctcatga cagcaagttc aatgtttttg ccacttgact gaaccacttc caggagtgcc 240  
 ttgataccca gcttaatggt cagatcatct gcttcaatgg ctccgltcagl atagttcttc 300

t

301

<210> 235  
 <211> 283  
 <212> DNA  
 <213> Homo sapien

<400> 235

tggggctgtg catcaggcgg gtttgagaaa tattcaattc tcagcagaag ccagaatttg	60
aattccctca tcttctaggg aatcatttac caggtttggg gaggattcag acagctcagg	120
tgctttcaat aatgtctctg aacttctgtc cctctttgtt catggatagt ccataaata	180
atgttatctt tgaactgatg ctcataggag agaataaag aactctgagt gatatonaca	240
ttagggattc aagaaatat tagatttaag ctccactgg tca	283

<210> 236  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 236

aggtcctcca ccaactgcct gaagcacggt taaaattggg aagaagtata gtgtagcata	60
aatactttta aatcgatcag atttccctaa ccacatgca atcttcttca ccagaagagg	120
tgggagcagc atcuttaata ccaagcagaa tgcgtaatag tcaaatataa tggatatatg	180
tgggtagacg gtttcatgag taccagtgtac tgggtatcag taatctggac ttgggttgta	240
aagcatcgtg taccagtcag aaagcatcna tactcgacat gaacgaatat aagaacacc	300
a	301

<210> 237  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 237

cagtggtagt ggtggtggac gtggcgctgg tctggtgccc ttttttgggt ccggtcacaa	60
actcaatttt tgttcgctcc tttttggcct ttccaattt gteratctca attttctggg	120
ccttggtcaa tgcctcatag taggagtcct cagaacagcc atggggatca aacatatect	180
ttgggtagt ggtgccaaagc tcttcaatgg cacagaatgg atcagcttct cgtaaatcta	240
gggttcagaa attctttctt cctttggata atgtagtcca tatccattcc ctcttttate	300
t	301

<210> 238  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 238

ggagcaggttt tttttttttt ttttttgatg gtgcagaccc ttgctttatt tgtctgactt	60
gttcacagtt cagccccctg ctccagaaac caacgggccg gctaaggaga ggaggaggca	120
ccttgagact tccggagtcg aggtcttcca gggttcccca gccatcaat cttttctctg	180
acccccctgc tgggaagcag ctccctgggg ggtgggaatg ggtgactaga agggatttca	240
gtgtgggacc cagggtctgt tccrcacagt agggagtgga agggatgact aatttcttta	300
t	301

<210> 239  
 <211> 239



&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 239

ataagcagct agggatttct ttatttagta atgtcctaac ataaaagttc acataactgc	60
ttctgtttaa ccatgatact gagctttgtg acacccaga aataactaag agaaggcnaa	120
cataatcct tagagatcaa gaaacattta cacagttcaa ctgttcaaaa atagctcacc	180
attcagccag tsagtagagt gtgaatgcca gcatacacag tatacaggtc cttcaygga	239

&lt;210&gt; 240

&lt;211&gt; 300

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 240

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gggatctgcc ctccagtggg accttttbaag gaagaggtgg gcccaagcta agtcccat	120
gctgggtgag ccagatgact tctgttccct ggtcaccttc ttcaatgggg cgaatggggg	180
ctgccaggtt tttaaaatca tgccttcact tgagacacac ggtcccttca cctcctcac	240
gctgtgggtg tcttttgatg aaaataccca ctttgttggc cttctgaag ctataatgtc	300

&lt;210&gt; 241

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 241

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cctcttttga ggaacttcca gcagctatgt tgggtctctt gagggaatgc aacaggtctg	120
ctcctccatg tattggaaaa ctgcaactg gactraactg gaaggagtg ctgctgccag	180
tgtgagaaac cagcctgagg tgacagaaac ggaagcaaac aggaacagcc agtctttct	240
tctctctct gtcatacggc ctctctcaag cttctttctg tgtcaggggc ctaaaaggga	300
g	301

&lt;210&gt; 242

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 242

ccagaggtct gggatgcac caatcactct gtttcacgtg acttttatca ccatacaatt	60
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gtcttcaaga atatatcatt cctttttcac tgaacccat tcaaatata agtcaagaac	180
cttaatatca acaatatat caagcaaac ggaaggcaga ataactacca taatttagta	240
taagtaccca aagttttata aatcaaaaag cctantgata accattttta gaattcaatc	300
a	301

&lt;210&gt; 243

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 243

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tgacgtgcag tcggactctg tggcccaagg gtatggctct ctgggcatga tgaccagcgt 180  
gctggtttgt ccagatggca agacagtga agcagaggct gccacggga ctgtaacccg 240  
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t 301

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<212> DNA  
<213> Homo sapien

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ccagggaacct tgganacagt tgacactgta aggtgcttgc tccccaagac acatccctaa 180  
agggtgttgt atggtagaaa cgtcttccct ctttatttgc cctcttatt tatgtgaaca 240  
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<211> 301  
<212> DNA  
<213> Homo sapien

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aaggccagga gatattgtca ttaatgtara ctccaggaca cttagagtata gcagccctat 180  
gttttcaaa agcagagatg caattaaata ttgtttagca tcaaaaaggc caatcaatac 240  
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g 301

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<212> DNA  
<213> Homo sapien

<400> 246  
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agtgccttct gtgaanatta aataaaacag ttaattcaaa gccttgatat atgttaccac 180  
taacaatcat actaatata ttttgaagta caaagtttga catgctctaa agtgacaacc 240  
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<212> DNA  
<213> Homo sapien

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gtgtcctgtg ttccgggtgc acacacaatc ctcattggga caggatcac catgcctgc 180  
ccttgatgat caagattggg gctcaagtgg attaagggag gcaagttctg ggttcttgc 240  
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a 301

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 gtacatttcc gctgtttggc aartccataa aacatttca gattttaate ccgaatttag 240  
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 c 301

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 <212> DNA  
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 ccagggagac acagcagtgä ctacagagctg gtcgcacact gtgctcctt cctcaccgac 180  
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 cataagcaca ccagctacttt tctctggctg gaatagttaa cttaagtatg gtacatctac 180  
 ctaaaagact actatgtgga ataatacata ctaatgaugt attacatgat ttacagacta 240  
 caataaaacc aacatgtctt ataacattaa gaaaaacaat aaagatacat gattgaaacc 300  
 a 301

<210> 251  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 251  
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 gccaggggtc ctcaaaaatg ccactgtcac tgcaggaaä tcttcttgag cagtacacct 180  
 cattgggac aatgaaaagc ttcaagaaat ctccaggtc actctcttga aggcceggaa 240  
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 c 301

<210> 252  
 <211> 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 252

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ctttctacat tctagaatca agagtgttaa taaatgtata tctatgtctt caagaatata	120
ccattctttt ttcactagga acccattcaa catataagtc aagaatctta atatcaacaa	180
atataatcag caacttggaa ggcagaataa ctaccataat ctagtataag taccacaaagt	240
tttatsaatc aaaagcccta atgataacra tttttagaat tcaatcatca ctgtagaatc	300

a

302

&lt;210&gt; 253

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 253

tcccttaaga agatgttact ttgttgggtt ttgttccccc tccatctcga ttctcgtacc	60
caactaanaa aaaaaataa agaaaaaatg tgcctgcgtc tgaanaataa ctctcttagct	120
tggtctgatt gtcttcagac cttaaaatat aaacttgttt cacaagcttt aatccatgtg	180
gatttttttt cttagagAAC cacaacaacat aaaaggagca agtcggactg aatacctgtt	240
tccatagtgc ccacagggta ttcctcacat ttctccata ggaaaatgct tcttcccaag	300

g

301

&lt;210&gt; 254

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 254

cgtcgcgct ttccttggg ggaagggcraa ggcagaggg ggtccaaagt cagcacgagg	60
aacttgacca attccttga agcgggtgg ttaaacctg taaatggga caaatccgc	120
craatctct tcatcttacc ctggtagact cctgactgta gaattttttg gttgaaacaa	180
gaaaaaata agcttttgg cttttcaag ttgttaaca ggtactgaa gactggcctc	240
acttaaatg agccaggaaa agctgcagat ttattaatg gctgttagt gtgcagtgc	300

c

302

&lt;210&gt; 255

&lt;211&gt; 302

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 255

agctttttct ttttttttt ttttttttt ttcattaaaa aatagtgtc tttattataa	60
attactgaan tgttttttt ctgaatataa atataaatat gtgcaaagt tgaacttggat	120
tgggattttg ttgagttctt caagcatctc ctataccct caagggcctg agtagggggg	180
aggaaaaagg actggagggtg gaatcttctt aaaaaacaag agtgattgag gcagattgta	240
aacattatta aaaaacaaga aacaaacaaa aaatatagga aaaaaaacac cccaaacacac	300

aa

302

&lt;210&gt; 256

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 256

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gttccagaaa acattgaagg tggottccca aagtcctaact agggatarcc cctctagcct      60
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aggcaaatag ctgctggcaa actggcatca cctggcttctt ggggatgggg gggcaagtgc      240
gtggcctctc ggcctgggta graagaacat tcagggtagg cctaagttan tctgtttagt      300
t

```

&lt;210&gt; 257

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 257

```

gttctggagg aactctggct tgcctattaa gtccactga ttttcaactat cccctgaatt      60
tccccactta ttttctctt tcaatatcc aggccttaga agaggctctac ctgcctccag      120
tcttacctag tccagtctac cccctggagt tagaatggcc atcctgaagt gaaaagtaat      180
gtcacattac tcccttcagt gatttcttct agnagtgcc atccctgaat gccaccaaga      240
tcttaattct cactcttcta atcttatctc ttgactcct ctttacaacc gagaaggctc      300
c

```

&lt;210&gt; 258

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 258

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cagragtagt agatgccgta tgcagracg cccagcactc ccaggatcag caccagcacc      60
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cccagggcaa caagaatcca ataccaggac tgggcaaaat ctccaagat cttaacactg      180
atgtctcggg cattgaggct gtcaataana cgtgatccc ctgctgtatg gtggtgtcat      240
tggtgatccc tgggagcgcc ggtggagtaa cgttggctcc tggaaagcag cgccccaac      300
t

```

&lt;210&gt; 259

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 259

```

tcatatctgc aaacaaatgc agactangcc tcaggcagag actaaaggac atctcttggg    60
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gcaagcccat aggaagccc aggatccctt gtgatcagga agtgggcccag gaaggctctgt    180
tccagctcac atctcatctg catgcagcac ggaccgcatg cggccactgg gtcttggctt    240
ccctcccatc ttctcaagca gtgtcctctg tgagccattt gcctccttgg ctccaggtgg    300
c                                          301

```

&lt;210&gt; 260

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 260

```

ttttttttct cccaaaggaa aaagaaggaa caggtctcat aaaaccat atagcaatggg    60
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agaactgtaa cagccacagt tggccatttc atgcraatgg cagcaaacaa caggattaac    180
tagggcaaaa taaataagtg tgtggaagcc ctgataagtg cttaataaac agactgattc    240
actgagacat cagtacctgc cggggcgggc gctcgagccg aattctgcag atatccatca    300
c                                          301

```

&lt;210&gt; 261

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 261

```

aaatactcga gcaaatccctg taactaatgt gtctccataa aaggctttga actcagtgaa    60
tgtgcttcca tccacgattc tagcaatgac ctctoggaca tcaagctcc tcttaagggt    120
agraccaact attccatcca attcatcagc aggaatbaaa ggctcttcag aagggtccat    180
ggtgacatcc aatttcttct gataatttag attctcaca accctctcag ttaagtgaag    240
ggcatgatga tcatcnaag cccagtggtc acttactcca gactttctgc aatgaagatc    300
a                                          301

```

&lt;210&gt; 262

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 262

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gaggagagcc tgttacagca ttgttaagca cagaatactc caggagtatt tgtuattgtc    60
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gggctttctg gtgcacacct aattttgtgc atctttgccc taatcctgg attagtcccc    240
catcattacc cccacattat aatgggatag attcagagca gatactctcc agcaagaat    300
c                                          301

```

&lt;210&gt; 263

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 263

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taatgaotga	cttcccagta	aggctctcta	aggggttaagt	angaggatcc	acaggatttg	240
agatgctaag	gccccagaga	tcgtttgata	caacctctct	attttcagag	gggaaaatgg	300
g						301

&lt;210&gt; 264

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 264

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gtggatagat	ctagaattgt	aacattttca	gaaaaccata	acatttgaca	gatgggaaag	180
ctcaattata	gatgcgaagt	tataactaaa	ctactatagt	agtaaaggaa	tacattttcc	240
acctttcata	taatttcaat	atcttggctt	gagggaactcc	acaaaatgta	tcacgtgcct	300
a						301

&lt;210&gt; 265

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 265

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catattcttg	gaagtctcta	atcaactttt	gttccatttg	tttctttct	tcaggaggga	180
ttttcagttt	gtcaacatgt	tctctaaaca	cacttgccca	tttctgtaaa	gaaatccaaag	240
cagtcgaagg	ctttgacatg	tcaacaacca	gcataactag	agcctctctc	agagatacgg	300
c						301

&lt;210&gt; 266

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 266

taccgtcttg	ctttccctcc	atccaggcca	tctgcgaatc	taratgggtc	ctcttattcg	60
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atagagacac	caataccctat	aacctctctc	ctaagctctc	ctataaccca	gggtgcacag	240
cacagactcc	tgacaactgg	taaggccaat	gaactgggag	ctcacagctg	gctgtgcctg	300
a						301

&lt;210&gt; 267

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 267

aaagagcaca	ggcagctca	gcccgccttg	gccatctaga	ctcagcctgg	ctccatgggg	60
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gttttcagtg ctgagtcocat ccaggaaaag ctcacctaga ccttctgagg ctgaatcttc      120
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ctcatttga ttctctctct tcttttcttt caagttggct ttcttcacat cctctgttc      240
aattcgcttc agcttgctg ctttagccct cattccaga agcttcttct ctttggcacc      300
t                                                                                   301

```

&lt;210&gt; 268

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 268

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aatgtctcac tcaactaact cccagertac cgtggcctaa ttctgggagt tttcttctta      60
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tgctgggtgg ctacgtgagc ccttttggag aaagcaagta ttattcttaa ggagtaacca      240
cttcccatgg ttctacttcc taccatcatc aattgtatat catgtattct ttggagaact      300
a                                                                                   301

```

&lt;210&gt; 269

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 269

```

taacaatata cartagctat ctttttaact gttcatcatt agcaccaatg aagattcaat      60
aaaattacct ttattcacac atctaaaaac aattctgcaa attcttagtg aagttaact      120
atagtcacag accitaaata ttacatttgt ttctatgtc tactgaaat aagttcacta      180
ctttcttgga tattctttac aaatcttat taaaattcct ggtattatca cccccaatta      240
tacagtacga caaccacctt atgragtttt taccatgatag ctctgtagaa gtttcacatc      300
t                                                                                   301

```

&lt;210&gt; 270

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 270

```

cattgaagag cttttgcga accatcagaac acaagtgctt ataanattaa ttaaggcetta      60
cacaagaata catattcctt ttatttctaa ggagttaaac atagatgtag ctgatgtgga      120
gagcttgctg gtgcagtgc tattggataa cactattcat ggccganitg atcangtcaa      180
craartcctt gaactggatc atcagaagaa ggggtgtgca cgatatactg cartagataa      240
tggacraacc aactaaattc tctcaccagg ctgcatcagt aaactggctt aacagaaaac      300
a                                                                                   301

```

&lt;210&gt; 271

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G



&lt;400&gt; 271

```

aaaagggtct ctaagattca acaattttaa taantatttg ataggacatt ctttctcatt      60
tttatagetc atcttttagg ttgatattca gttcatgett cctttgctgt tcttgatcca      120
gaattgcaat caettcatca gcttgattc gctccaatto tctataaagt ggggtccaagg      180
tgaaccacag agccacagca cactctttc ccttggtgac tgccttcacc ccattganggt      240
tctctcctcc agatganaac tgatcatgcg ccacatttt ggggtttata gaagcagtc      300
c

```

&lt;210&gt; 272

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 272

```

taatttgctc agccacagat aacaccaatc aattggaaca atcactgtc ttcaaatgtc      60
ttatcagaa acccaatgag cttggaatct tcataatacc taacatgac gtatttagga      120
tcaataaatt cctcatgat gagraagaaa aattctttgc gcacccctc tgcaccaca      180
gcctcttctc caaccaatct aaccttgagt ggttctttgt aatctatgtt ctttgttttc      240
ctaaggactt ccattgcac cctacasta tttctctac gcaccactag aattagcag      300
g

```

&lt;210&gt; 273

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 273

```

aatgtgtgt atgtgtatct ttgggaanan aanaagacat cttgtttayt attttcttgg      60
agagangctg ggacatggt aatcacwtaa ttgtctayta tyactttaat ctgactygaa      120
gaaccgtcta acaataaaat ttaccatgtc dtatattcct tatagtatgc ttatttcacc      180
tcttctgtt ccagagagag tatcagtgc ananatttma gggcgaamac atymattgggt      240
gggacttnty ttacngagm auctgcccg agcgcctcg makengant crgcsananc      300
t

```

&lt;210&gt; 274

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (301)

&lt;223&gt; n = A,T,C or U

&lt;400&gt; 274

```

cttatatact ctttctcaga ggcaanagag gagatgggta atgtagacaa ttctttgagg      60
aacagtaaat gattattaga gagaangaat ggaccaagga gacagnaatt aacttgtaaa      120
tgattctctc tggaaatctga atgagatcaa gaggccagct ttagcttctg gaaaagtcca      180
tctagstatg gttgcattct cgtcttcttt tctgcagtag ataattgagg aaccgaaggc      240
aattgtgctt cttttgataa gaagctttct tggatcatatc aggaatttc aganaagtc      300

```

c

301

<210> 275  
<211> 301  
<212> DNA  
<213> Homo sapien  
  
<220>  
<221> misc\_feature  
<222> {1}...{301}  
<223> n = A,T,C or G

<400> 275

tgggtgtcag	cagcaagtgg	cattgaacat	tccaatgtgg	agcccaaac	acagaaaatg	60
gggtgaaatt	ggccaacttt	ctatcaactt	atgttggcua	ttttgccacc	aacagttaagc	120
tggcccttct	aataaaagaa	aattgaaagg	tttctoacta	aacgggatta	agtatgtggag	180
tcaagagact	cccagggctc	agcgtacutg	cccgggaggc	cgctcgaagg	cgaattctgc	240
agatcccat	cacactggcg	gncgctcgan	catgoatcta	gaaggaccaa	ttcgccttat	300
a						301

<210> 276  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 276

tgtacacata	ctcaactaat	aaatgactgc	attgtggtat	tattactata	ctgattacat	60
ttatcatgtg	acttctaatt	agaaatgta	tccaaagca	aaacagcaga	tatcacaaat	120
ttaagggaca	gaagataaac	actaacagat	aaggcaactt	atacatttag	aatccaaatc	180
caatacatit	aaacatttgg	gaatgagggg	ggacaaatgg	aagccagatc	aaatttgtgt	240
aaaactattc	agtatgttcc	ccttgcttca	tgtctgagaa	ggctctcctt	caatggggat	300
g						301

<210> 277  
<211> 301  
<212> DNA  
<213> Homo sapien

<220>

<221> misc\_feature  
<222> {1}...{301}  
<223> n = A,T,C or G

<400> 277

tttgttgatg	tcagtatttt	attacttgcg	ttatgagtgc	tcacctggga	aattctaaag	60
atacagagga	cttggaggga	gcagagcaac	tgaatttaat	ttaaaaggaag	gaatacattg	120
gaatcatggc	aotcctgata	ctttcccaa	tcaacactct	caatgcccc	ccctcgtcct	180
caacatagtg	gggagactaa	agtggccacg	gatttgcctt	angtgtgcag	tgcgttctga	240
gttctctgtc	gattacatct	gaccagtctc	ctttttccga	agtcctctcg	tccaattctg	300
c						301

<210> 278  
<211> 301  
<212> DNA  
<213> Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 278

taccactaca ctccagcctg ggcacacagag caagacctgt ctcaaaagcat aaaatgggaat	60
aacatatacaa atgaaacagg gaaaatgaag ctgcacattt atggaagcca gggcttgtca	120
cagtctctac tgttattatg cattacctgg gaatttatat aagcccttaa tcaaatgcc	180
aatgaacata tcatgtgtgc tcacaatggt ctggcactat tataagtgtc tcacaggttt	240
tatgtgttct tcgtaacttt atggantagg tctcggccg cgaacacgct aagccgaatt	300
c	301

&lt;210&gt; 279

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 279

aaagcaggaa tgacaaagct tgccttctctg gtatgttcta ggtgtattgt gacttttaact	60
gttatattaa ttgccaatat aagtaaacat agattatata tgtatagtgt ttcaaaaagc	120
ttagaccttt acctccagc caccacacag tgccttgatct ttcagagtea gtcatttgggt	180
atacatgtgt agttccaaag cacataagct agaaanaanaa atatttctag ggagcactac	240
catctgtttt cacatgaat gccacacaca tagaactcca acatcaattt catbgcacag	300
a	301

&lt;210&gt; 280

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 280

ggtactggag ttttctctcc ctgtgaaac gtaactactg ttgggagctga attgaggatg	60
tagaaagggt gtggaaccaa attgtggtca atggaatatg gagaatatgg ttctcaactct	120
tgagaaaaaa acctaaagatt agcccaggta gttgctgtga acttcagttt ttctgcctgg	180
gtttgatata gtttaggggt ggggttagat taagatctaa attacatcag gacaaagaga	240
cagactatta actccacagc taattaaagg ggtatgttcc atgtttattt gttaaagcag	300
t	301

&lt;210&gt; 281

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 281

aggtacaaga aggggaatgg gaaagagctg ctgctgtggc attgttcaac ttggatatcc	60
gccgagcaat ccaaatcctg aatgaagggg catcttctga aaaaggagat ctgaatctca	120
atgtggtagc aatggcttta tggggttata cggatgagaa gaactccctt tggagagaaa	180
tgtgtagcac actgcgatta cagctaaata acccgtattt gtgtgccatg ttgcatttc	240

tgacaagtga aacaggatct tacgatggag ttttgtatga aaacaaagtt gcagtaccta 300  
g 301

<210> 282  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 282  
caggctactac agaatataaa tactgacaag caagtagttt cttggcggtgc argaattgca 60  
tcragaaccc aaaaatttaag aatttcataa agacattttg tgggcacctg cttagcacaga 120  
agcgacagaag caaagcccag gcagaacctt gctaacttta cagctcager tgcacagaag 180  
cgcagaagca aagcccaggt agaacatgc taaccttaca gctcagcctg cacagaagcg 240  
cagaagcaca gccacggcag aacatgctta ccttacagct cagcctgac agaaagcag 300  
a 301

<210> 283  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 283  
atctgtatcc gccagacaaa ctttatarag tctagagagg tgagcgaag gatgcaaaag 60  
cacttrgagg gctttataat aatatgctgc ttgaaaaaaa aaatgtgtag ttgatactca 120  
gtgcacttcc agacatagta aggggttgc ctgaccaatc aggtgatcat tttttctatc 180  
acttcccag ttttatgcaa aaatttctgt aaattctata atggtgatat gaattcttta 240  
ggaacatat acatttctaa aaattctatt tatgtgaag ctgacagagc aatttgcctt 300  
g 301

<210> 284  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 284  
caggtaaaaa acgtatttaa gtggctcaga atttgaacat ttgtggtctt tcttctactt 60  
gcttctgtgt tggycaaaag aacatcttcc ctAAatatat attaccaaga aaagcaagaa 120  
gcagattagg tttttgacaa acaaaaacgg ccaaaagggg gctgacctgg agcagagcat 180  
ggtgagaggc aaggcatgag agggcaagtt tgtgtgtgac agatcctgtc ctactttatc 240  
actgggttaa aagaaaaaaa agttcattga tgtgaagga tatatacagt gttagaaatt 300  
a 301

<210> 285  
<211> 301  
<212> DNA  
<213> Homo sapien

<220>  
<221> misc\_feature  
<222> 11}... (301)  
<223> n = A,T,C or G

<400> 285  
acatcaccat gatcggatcc cccaccatt atacgttcta tttttacata aatattcttc 60  
aatgatcat agtgtttta aaaaaatacc gaaaacttct tctgcatccc aatctctaac 120

```

caggaaagca aatgctattt acagaactgc aagccctccc tcaaacnana ctatttctgg 180
attaaatag tctgacttct tttgaggtca cagactagg caaatgctat ttacgatctg 240
caaaagctgt ttgaagagtc aaagccccc tgtgaacacg atttctggac cctgtaacag 300
t 301

```

```

<210> 286
<211> 301
<212> DNA
<213> Homo sapien

```

```

<400> 286
taccactgca tccagcctg ggtgacagag tgagactccy tctccaaaa aacttttctg 60
tgtatattat tttgacctta cagtggatca ttctagttag aaggyacagt aagatttttt 120
atcaaaatgt gtcctgccag taagagatgt tatattcttt tctcatttct tccccaccca 180
aaaataagct accatatagc ttataagctc caattttttg ccttttacta aaatgtgatt 240
gtttctggtc attgtgtatg ctccatcacc tatattagga aaattccatt ttttcccttg 300
t 301

```

```

<210> 287
<211> 301
<212> DNA
<213> Homo sapien

```

```

<400> 287
tacagatctg ggaactaaat attaaaaatg agtgtsacctg gatetatgga gaactgttggg 60
cccagaagga acgtagagat cagatattac aacagcttly ttttgagggg tagaatatg 120
aaatgatttg gttatgaacg cacagttag gcacagaggr cagaatctg accctctgcc 180
cgtggttat ctctctccca gtttggctgc ctcatgttc cacagtattc catttctgtt 240
gttgcatgic ttgtgaagcc atcaagattt tctcgtctgt tttctctcca ttggbaatgc 300
t 301

```

```

<210> 288
<211> 301
<212> DNA
<213> Homo sapien

```

```

<400> 288
gtacacctaa ctgcaaggac agctgaggaa tgtatcgggc agccgctttt aaagaagtag 60
agtcaatagg aagacaaatt ccagttccag ctcagtctgg gtatctgcaa agctgcaaaa 120
gatcttttaa gacaatttca agagatatt tctttaaagt tggcaatttg gagatcatac 180
aaaagcatct gcttttgcga tttaatttag ctcatctggc cactgggaaga atccaaacag 240
tctgccttaa ttttggatga atgcattgat gaaattcaat aatttagana gttaaaaaaa 300
a 301

```

```

<210> 289
<211> 301
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (301)
<223> n - A,T,C or G

```

```

<400> 289

```

```

ggtagactgt ttcacgttta tgtttctaca cattgctacc ttagtgctcc tggaaactta      60
gtttttgatg tctccaagta gteraccttc atttacctct ttgaaactgt atcactcttg      120
craagtaaga gtgggtggct atttcagctg ctttgacaaa atgactggct cctgacttaa      180
cgttctataa atgaatgtgc tgaagcaaa tgccaatggg ggccggcgaan aagagaaaga      240
tgtgttttgt tttggactct ctgtggtccc ttcacatgct gtgggtttcc aaccagngga      300
a                                          301

```

&lt;210&gt; 290

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 290

```

aacctgggct cttcttgata aatataraga atgcttggca tatcacaagat tctatactac      60
tgactgatct gttcatttct ctacagctc ttaccccaa aagcttttcc accctaagtg      120
ctctgacctc cttttctaatt cacagtaggg atagaggcag anccacctac aatgaacatg      180
gagttctatc aagaggcaga aacagcacag atccccagtc ttaccattcg ctagcagtgc      240
tgccttgaac aaaaacattt ctocatgtct cabtttcttc atgcttcaag taacagtgag      300
a                                          301

```

&lt;210&gt; 291

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 291

```

caggtaccac ttctttctat cctagaaacn tttcatttta tgttgttgaa acataacaac      60
tatatcagct agatttttct tctatgttct acctgctatg gaaaacttga cacattctgc      120
tttactcttt tgtttatagg tgaatcaca aatgtatttc tatgtattct gtagtctaat      180
agccatggct gtttacttca ttttaattat ttagcatcaa gacattatga aaaggcctaa      240
acatgagctt caatttccca ctacctaatt agcatctgtt atttcttaac cgtaatgcct      300
a                                          301

```

&lt;210&gt; 292

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}...(301)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 292

```

accttttagt agtaatgtct aataataaac aagaaatcaa ttttataagg tccatatagc      60
tgtattaat aactttcaag ttttaaagat aaaataccat catttcaant gtgggtatct      120
aaaaccnaag natataaccg aaaggaaaaa cagatgagac ataaaatgat ttgcagatg      180
ggaaatatag tacttyatga atgttnatta aattocagtt ataatagtgg ctacacactc      240
tcactacaca cacagacccc acagtcttat atgccacaaa cacatttcca taacttgaaa      300
a                                          301

```

<210> 293  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 293  
 ggtaccaagt gctggtgcca gccgtttacc tgtttctcact gaagaagtctg gctaattgctc 60  
 ttgtgtagtc actttctgatt ctgacaarca atcaatcaat ggcctagagc actgactgtt 120  
 aacacaaaag tcaactagcaa agtagcaaca gctttaagtc taatacaaa gctgttctgt 180  
 gtgagaattt tttaaaaggc tacttgata ataacccttg tcatTTTTTaa tgaactcgg 240  
 ccgcgaccac gctaagcaga attctgcaga tctccatcac actggcgggc gctcgagcat 300  
 g 301

<210> 294  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (301)  
 <223> n - A, T, C or G

<400> 294  
 tgaccataa caatatacac tagctatctt cttaactgtc catcattagc accaatgaag 60  
 attcaataaa attaccttta ttcacacac tcaaaacaa tctgcaatt cttagtgaag 120  
 tttactata gtacaganc tcaactatc acattgttt ctatgtctac tgaataaag 180  
 ttcactaatt ttctgggata ttctttacaa aatcttatta aaatccctgg tattatcacc 240  
 ccaattata cagtagcaca accacettat gtagtlttta catgatagct ctgtagaggt 300  
 t 301

<210> 295  
 <211> 305  
 <212> DNA  
 <213> Homo sapien

<400> 295  
 gtacttttc tctccctcc tctgaattta attctttcaa ctgcgaattt gcaaggatta 60  
 cacatttcac tgtgatgtat attgtgtgtc aaaaaaaa gtgtctttgt ttaaaattac 120  
 ttggtttgtg aatccatctt gctttttccc cattggaact agtcattaac ccattcttga 180  
 actggtagaa aaactcttga agagctagtc taccagcacc tgacaggaga attggatggg 240  
 tctcagaacc atttcaccca gacagcctgt tcttatcctg ttaataaaat tagtttgggt 300  
 tctct 305

<210> 296  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 296  
 aggtactatg ggaagctgct aaataatat ttgatagtaa aagtatgtaa tgtgctatct 60  
 caactagtag taactaaaa ataaactgaa actctatgga ctctgaagtt atttccctg 120  
 attaataga attaataaac caatatgagg aaecatgaaa ccatgcaatc cactatcaac 180  
 tttagaaaag tgattgaacg aaccacttag ctttcagatg atgaacaactg ataagtcatt 240

tgctactact ataaatttta aaattctgta ataagatgga ctacagggag gaaaaagggg 300  
c 301

<210> 297  
<211> 300  
<212> DNA  
<213> Homo sapien  
  
<220>  
<221> misc\_feature  
<222> (1)...(300)  
<223> n = A,T,C or G

<400> 297  
actgagtttt aactggacgc caagcaggca aggttgaag gtlttgctct ctttgtgcta 60  
aaggttttga aaaccttgaa ggagaaatrat ttgacaaga agtacttaag agtotagaga 120  
acaaagangt gaaccagctg aaagctctcg ggggaanctt acatgtgttg ttaggcctgt 180  
tccatcattg ggagtgcact ggcaatccct caaaatttgt ctgggctggc ctgagtggtc 240  
accgcacctc ggccgcgacc acgctaagcc gaattctgca gatatccatc acactggagg 300

<210> 298  
<211> 301  
<212> DNA  
<213> Homo sapien  
  
<220>  
<221> misc\_feature  
<222> (1)...(301)  
<223> n = A,T,C or G

<400> 298  
tatggggttt gtcaacccaa agctgatgct gagaagggcc tccctggggc ccttcccgag 60  
ggcatctgag agacctggg ttccagtgtt tctggaaatg ggtccuagtg ccgccggctg 120  
tgaagctctc agatcaatca cgggaagggg ctggcggtgg tggccacctg gaaccacctt 180  
gtcctgtctg ttacalttc actaycaggt ctctcttggg cattacnati tgttccctta 240  
caacagtgac ctgtgcattc tgcgttggcc tgcgtgtgct gcaggtgget ctacgcgagg 300  
t 301

<210> 299  
<211> 301  
<212> DNA  
<213> Homo sapien

<400> 299  
gttttgagac ggagttttcac tcttgttgco cagactggac tgcraatggca gggtrctctgc 60  
tcaactgcacc ctctgcctcc caggttcgag caattctccr ggcctagcct cccaggtagc 120  
tgggatttga ggtcacgac accataccra gctaattttt ttgtattttt agtagagacg 180  
gagtttcgac atgtcggcca gctggcttca aartccctgac ctcaagcgac ctgcttgcct 240  
cggcttccca aagtgttggg attataggca tgagtcaaca cgtccagcct aaagatattt 300  
c 301

<210> 300  
<211> 301  
<212> DNA  
<213> Homo sapien



<400> 300  
 attcagtttt atttgcctgc ccagtatctg taaccaggag tggcacaas tcttgccaga 60  
 tatgtccac acccaactgg aaaggctccc acctggctac ttctctatc agcvgggtca 120  
 gctgcattcc acaagggttct cagcctaatt agtttcarta cctgccagtc tcaaaactta 180  
 gtaagcaag accatgacat tccccacgg aaatcagagt ttgccccacc gtcttggtac 240  
 tataaagcct gctctaacca gtcttgctt ctccacacca atcccgagcg catcccccat 300  
 g 301

<210> 301  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 301  
 ttaaatTTTT gagaggataa aaaggacaaa taatctagaa atgtgtcttc ttcagtcctgc 60  
 agaggacccc aggtctccaa gcaaccacat ggtcaaggyc atgaataatt aaggttggc 120  
 ggggaactc ac aagacccctc agagctgaga ccccccaac agtgggagct cacaaagacc 180  
 cttagagctg agacacccc aacagcggga gctcacaaag accctcagag ctgagacacc 240  
 caaacagca cctcgttcag ctgcccacat gctgaataag gatgcaatgt ccagaagtgt 300  
 c 301

<210> 302  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 302  
 aggtacacat ttagcttctg gtaaatgact cccaaacchg atttcaaaat caagttaatg 60  
 tgaattttg aaattactac ttaattctaa ttcaaataa caatggcatt aaggtttgac 120  
 ttgagttggc tcttagctat atttatggta aataggctct taccacttgc aataaactgg 180  
 ccacatcatc aatgactgac ttccctgtaa ggctctctaa ggggttaagta ggaggatcca 240  
 caggatttga gatgctaagg cccagagat cgtttgatcc aacccctctta ttttcagagg 300  
 g 301

<210> 303  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 303  
 aggtaccac tggggaataa ggtagaggat cttttttct tcccatatca actaagttgt 60  
 atattgtct ttgacagttt aacacatct ctctgtcag agattctttt acaatagcac 120  
 tggctaattg aactacrgct tgcattgtaa aaatggctgt ttgtgaaatg atcataggcc 180  
 agtaacgggt atgttttct aactgatctt ttgtctgtc caaagggacc tcaagacttc 240  
 catcgattt atattctggg tctagaaaag gaggtaatct gttttccctc ataaattcac 300  
 c 301

<210> 304  
 <211> 301  
 <212> DNA  
 <213> Homo sapien

<400> 304  
 acatggatgt tatcttgag actgtcacc tgaatttgta ttgcttgac atggccta 60

tattagcttc agtttcagct taccracctt ttgtctgca catgcaraab agacagtgc	120
cttttttagtg tatcatatca ggaatcatct cacattgggt tgtgccatta ctggtgcagt	180
gactttcagc cacttgggtt aggtggagtt ggcacatgt ctccactgca aattactga	240
tttcccttt gtaattaata agtgtgtgtg tgaagattct ttgagatgag gtatatatct	300
c	301

<210> 305

<211> 301

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(301)

<223> n, - A,T,C or G

<400> 305

gungtacagc gtggtcaagg tacaagaag aaaaaaatgt gagtggcatc ctgggatgag	60
cagggggaca gacctggaca gacacgttgt catttctgc tgtgggtagg aaatgggag	120
ttaaggaggc gaaacagata cnaaatctcc auctcagtt taaggtattc tcatgctag	180
aatattggtt gaacaaaga tacattcata tggcaataa ctaaccatgg tggaaacaaa	240
ttctgggatt taagttggat accaangaaa ttgtattaa agagctgttc atggaataag	300
a	301

<210> 306

<211> 8

<212> PRT

<213> Homo sapien

<400> 306

Val Leu Gly Trp Val Ala Glu Leu

1

5

<210> 307

<211> 637

<212> DNA

<213> Homo sapien

<400> 307

acaggggatg aagggaaagg gagaggatga ggaagcctcc ctggggattt ggtttggtcc	60
ttgtgatcag gtggtctatg gggcttatcc ctacaaagaa gaatccagaa ataggggcac	120
attgaggnat gatacttgag cccaaagagc attcaatcat tgttttattt gccttmtttt	180
cacacccttg gtgagggagg gattaccacc ctggggttat gaagatgggt gaacacccca	240
cacatagcac cggagatatg agatcaacag ttcttttagc atagagattc acagcccaga	300
gcaggaggac gcttgcacac catgcaggat gacatggggg atgggctcgg gattgggtgtg	360
aagaagcaag gactgttaga ggcaggcttt atagtaacaa gacgggtggg caaactctga	420
tttccgtggg ggaatgtcat ggtcttctt tactaagttt tgagactggc aggtagttaa	480
actcattagg ctgagaacct tgtggaatgc acttgaccca scgatagag gaagtagcca	540
ggtgggagcc ttccccagtg ggtgtgggac atatctggca agattttgtg gcaactcttg	600
ctacagatac tggggcagca aataaactg aactctt	637

<210> 308

<211> 647

<212> DNA

<213> Homo sapien

&lt;220&gt;

&lt;221&gt; Misc\_feature

&lt;222&gt; (1) ... (647)

&lt;223&gt; G - A,T,C or G

&lt;400&gt; 308

acgattttca	ttatcagcta	aatcggttca	ctcaaggggc	caaccacagc	tgggagccac	60
tgttcagggg	aagggttata	tgggactttc	tactgoccaa	ggttctatac	aggatataaa	120
ggngcctcac	agtatagatc	tggtagcaaa	gaagaagaaa	caaacactga	tctctttctg	180
ccacccctct	gaccccttgg	aactcctctg	cccctttaga	acagagcctac	ctaatactctg	240
ctagagaaaa	gaccaacaaac	ggcctcaaaag	gatctcttac	catgaagggtc	tcaactaatt	300
cttggttaag	atgtgggttc	cacattaggt	tctgaatatg	gggggaagggt	tcaatttgcct	360
catttttgtgt	gtggataaaag	tcaggatgcc	cagggggcag	agcagggggc	tgtttgtctt	420
gggaacacctg	gotgagcata	taaccatagg	ttatggggag	caaacacaaca	tcaaatgtac	480
tgtatcaatt	gccatyaaga	cttgaggggc	ctgaatctac	cgattcatct	taaggcagca	540
ggacragttc	gagtggcaac	aatgcagcag	cagaatcaat	ggaaacaaca	gaatgattgc	600
aatgtccttt	ttttctctct	gcttctgaat	tgataaaagg	ggacct		647

&lt;210&gt; 309

&lt;211&gt; 460

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 309

actttatagt	ttaggctgga	cattggaaax	aaaaaaagc	cagaacacaa	tgtgatagat	60
aatatgatcg	gttgcacart	tcagactga	tgaatgatga	acgtgatgga	ctattgtatg	120
gagcacatct	tcagaaagag	ggggaaatac	tcatactttt	tggccagcag	ttgtttgatc	180
accaaacatc	atgocagaat	actcagcaaa	ctttcttagc	tcttgagaag	tcaaatgccg	240
ggggaaattta	ttcttggcaa	ttttaatttg	actccttatg	tgagagcagc	ggctacccag	300
ctgggggtgg	ggagcgaacc	cgtcactagt	ggacatgcag	tggcagagct	cctggttaac	360
acctagaggga	atacacaggc	aratgtgtga	tgccaagcgt	gacacctgta	gcactcaaat	420
ttgtcttggc	tttgtctttc	ggtgtgtaag	attctcaagt			460

&lt;210&gt; 310

&lt;211&gt; 539

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 310

acgggactta	tcaaataaag	ataggaaag	aagaaaactc	aatattata	ggcagaantg	60
ctaaagggtt	taaaatatgt	caggattgga	aggaaggcatg	gataaagaac	aaagtccagt	120
taggaaagag	aaacacagaa	ggaagagaca	caataaaagt	cattatgtat	tctgtgagaa	180
gtcagacagt	aagatttctg	ggaaatgggt	tggtttcttg	tatggtatgt	attttagcaa	240
taatctttat	ggcagagaaa	gtcaaaatcc	tttagcttgc	gtgaatgatc	acttgcctgaa	300
ttcctcaagg	taggcattgt	gaaggagggt	ttagaggaga	caragacaca	atgaactgac	360
ctagatagaa	agccttagta	tactcagcta	ggaatagtga	ttctgagggc	acactgtgac	420
atgattatgt	cattacatgt	atggtagtga	tggggatgat	aggaagggaag	aacttatggc	480
atattttcac	cccccacaaa	gtcagttaaa	tattggggaca	ctaaccatcc	aggtcaaga	539

&lt;210&gt; 311

&lt;211&gt; 526

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(526)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 311

caaatttgag ccaatgacat agaattttac aaatcaagaa gcttattctg gggccatttc	60
ttttgacgtt ttctctaaac tactaaagag gcattaatga cccataaatt atattatcta	120
catttacagc atttaaatg tgttcagcat gaaatattag ctacagggga agctaaataa	180
attaacatg gaataaagat ttgtccttaa atataatcta caagaagact ttgatatttg	240
tttttcacaa gtgaagcatt ctatataagt gtacataacct ttttggggaa aotatgggaa	300
aaaatgggga aactctgaag ggttttcaagt atcttacctg aagctacaga cttcataacc	360
tctctttaca gggagctcct gtagccctta cagaatgag tggctgagat tcttgattgc	420
acagcaagag cttctctctt aaacctttt cctttttagt atctgtgcat caagtataaa	480
agttctataa actgtagtnt acttatttta atccccaaag cacagt	526

&lt;210&gt; 312

&lt;211&gt; 500

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(500)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 312

cctctctctc cccacccctt gactctagag aaccggggtt tctccragta ctcagcaat	60
tcattttctga aagcagttga gccactttat tccaaagtac actgcagatg ttcaaaactct	120
ccattttctc ttcccttcca cctgcccagt ttgtctgact tcaacttgtc atgagtgtaa	180
gcatttaagga catttatgct ctctgattct gaagacaggc cctgctcatg gatgactctg	240
gcttcttagg acaatatttt tcttccaaaa ttagtaggaa ctctaaactt atccctcttt	300
tcagatgttc tagcagcttc agacatttgg ttangaaccc atggggaaaaa acaaaactct	360
tgctaattgc gtttcttttg taaccanga ttcttatttg nctggtatag aatctcagct	420
ctgaagtggt ggttaagatt ctgtgtttg antataggag aaatcagttt gctgaaaagt	480
tagtcttaat tatctatcgg	500

&lt;210&gt; 313

&lt;211&gt; 718

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(718)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 313

ggagatttct gtggtttcca gccaggggag accaggaaga totgcatggt ggganaggacc	60
tgatgataca ggggtgagaa ataagaaagg ctgctgactt taccatctga ggcacacat	120
ctgctgaat ggagataatt aacattracta gaaacagcaa gatgacata taatgtctaa	180
gtagtacat gtttttgcac atttccayer cttttaaata tccacacaca cagggaagcac	240
aaaagggaag acagagatcc ctgggagaaa tgcctggcgg ccatcttggg tcatcgatga	300
gctcgccct gtgcttcttc ccgtcttga ggganaggaa ttagaaaatg aattgatgtg	360
ttccttaag gatggcagga aaacagatcc tgttctggat atttatttga aagggtattc	420

agatttgaaa tgaagtcaca aagtgagrat taccoatgag aggaaaacag acgugaaaat	480
cttgatggtr cacaagacat gcaacaaaca aaatgggata ctgtgatgac acgagcagcc	540
aaetggggag gagataccac ggggtagagg tcaggyattct ggcctgctg cctaaactgtg	600
cgttatacca atcatttcta ttctaccct caaacaagct gtngaatatc tgacttacgg	660
ttcttntggc cccatcttc atnatccacc cctctctttt aannttctc caaantgt	718

&lt;210&gt; 314

&lt;211&gt; 358

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 314

gtttattttac attacagaaa aaacatcacg acnatgtata ctatttcaca tatatccata	60
cataatcana tatagctgta gtacatgttt tcattgggtg agattaccac aaatgcagg	120
caacatgtgt agatctcttg tcttattctt ttgtctataa tactgtattg tctagtcaca	180
gctctcggtg gtccagccac tgtgaaacat gctcccttta gattaacctc gtggagctc	240
ttgttgtatt gctgaactgt agtgcctgt attttgcttc tgtctgtgaa ttctgttct	300
tctggggcat ttccttgtga tgcagaggac caccacacag atgacagcaa tctgaatt	358

&lt;210&gt; 315

&lt;211&gt; 341

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 315

taccacctcc ccgctggcac tgatgagccg catcaccaatg gtcaccagca ccatgaaggc	60
ataggtgatg atgaggacat ggaatgggac cccaaggatg gctgtccaa agagcgagt	120
gaccccatc ctgaagatgt ctggaacctc taccagcagg atgatgatag cccaatgac	180
agtcaccagc tcccagacca gccggataac gtccctaggg gtcatgtagg ctccctgaag	240
tagcttctgc tgaagaggg tgttgtcccg ggggctctg cggctattgg tctgggctt	300
gagggggcgg tagatgcagc acatggctga gcgatgatg t	341

&lt;210&gt; 316

&lt;211&gt; 151

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 316

agactgggca agactcttac gcccacact graatttggc ctgtgtgccc tatccattta	60
tgtgggctt tctcagttt ctgattataa acaccactgg agcgatgtgt tgactggact	120
catcaggga gctctggtt caatatctag t	151

&lt;210&gt; 317

&lt;211&gt; 151

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 317

agaactagtg gatcctaag aaataactga aacatatatt ggcatttatc aatggctcaa	60
atcttcaatt atctctggcc ttaacctgg cctctgaggc tgcggccagc agatcccagg	120
ccagggtctt gttcttgcca caactgctt a	151

&lt;210&gt; 318

&lt;211&gt; 151

&lt;212&gt; DNA

<213> Homo sapien

<400> 318

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actggtggga ggcgtgttct agttggctgt ttccagaggg gtctttcggg gggacctctt    60
gttcgagggc ggagtgcttt tattcttggc gggagaccgc acattccact gctgaggctg    120
tggggggcgt ttatcaggca gtgataaaca t                                     151
```

<210> 319

<211> 151

<212> DNA

<213> Homo sapien

<400> 319

```
aactagtggg tccagagcta taggtacagt gtgatctcag ctctgcaaac acattttcta    60
catagatagt actaggtatt aatagataty taagagaaga aatcacacca ttaataatgg    120
taagattggg ttatgtgat tttagtgggt a                                     151
```

<210> 320

<211> 150

<212> DNA

<213> Homo sapien

<400> 320

```
aactagtggg tccactagtc cagtgtgggt gaattccatt gtgttggggg tctagatcgc    60
gagcpgctgc cctttttttt tttttttttg ggggggaatt tttttttttt aatagttatt    120
gagtgttcta cagttacag taaataccat                                     150
```

<210> 321

<211> 151

<212> DNA

<213> Homo sapien

<400> 321

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agcaactttt tttttcatcc aggttatctt aggccttagg tttctcttca cactgcagtt    60
taggttggca ttgtaccag ctatggcata ggtgttaacc aaaggctgag taacatggg    120
tgctctgag aatcaaat cticatatcc t                                     151
```

<210> 322

<211> 151

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(151)

<223> n = A,T,C or G

<400> 322

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atcagccttc ttctctgtt tcttgccttc ctctttcttc ttcttasatt ctgcttgagg    60
tttgggcttg gtcagtttg cacagggtt ggagatggg acagtcttct ggcattcggc    120
attgtcagg gtcgcttca nacttccagt t                                     151
```

<210> 323

<211> 151

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(251)

<223> n = A,T,C or G

<400> 323

tgaggacttg tktttttttt ctttattttt aatctcttta ckttgtaaatt atattgocct	60
nagactrant tactnccag ttgttggttt twtgggagaa atgtacttg acagttagct	120
gttcaatya aaagacactt anccatgtg g	151

<210> 324

<211> 461

<212> DNA

<213> Homo sapien

<220>

<221> misc\_feature

<222> (1)...(461)

<223> n = A,T,C or G

<400> 324

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agaagtgttc agctaaagga atccaggttg ttggttggac tgttaatacc ttgatgaaa	120
aggttacta cgaatcccat cttggttcca gctatctcac tgacagcatg gtagaagact	180
gcgaacctca cttctagact ttacaggttg gacgaacagg gttagaagac tgccaggggc	240
ctcatcaggg gatctcaaaa taccctttgt gctaccaggg ccttggggaa ttaggtgact	300
cacacaaatg caatagtttg ccactgcatt ttacactgaa ccaagctaa acccggtgtt	360
gccaccatgc accatggcat gccagagttc aacactgttg ctcttgaana ttgggtctga	420
aaabacgtac aagagccctt gacctgacct agctganga c	461

<210> 325

<211> 400

<212> DNA

<213> Homo sapien

<400> 325

acactgtttc catgttatgt ttctacacat tgctacctca gtgtccctgg aaacttagct	60
tttgatgtct ccaagtagtc cacttccatt taactctttg aaactgtatc atctttgcc	120
agtaagagtg gtggcctatt tcaagtgttt tgacaaaatg actggctcct gacttaacgt	180
ctataaatg aatgttgtga agcaaatgct ccatgtgtgg ggcgaagaag agaaagatgt	240
gttttgtttt ggaactctctg tggtrccctt caatgctgtg ggtttccaac caggggaagg	300
gtcccttttg ctttgcraag tgcataaacc atgagcacta cgtaccatg gttctgcctc	360
ctggccaaag aggtgtgttt gcaagaatga aatgaatgat	400

<210> 326

<211> 1215

<212> DNA

<213> Homo sapien

<400> 326

ggaggactgc agnccgcact cgcagccctg gcaggcggca ctggtcatgg aaacgaatt	60
gttctgtctg ggcgtcttg tgcattccga gtgggtgtctg tcagccgcac actgtttcca	120
gaactcttac accatcgggc tgggcttgca cagctcttgag gccgaccaag agccagggag	180

ccaghtggtg	gaggccagcc	tctccgtacg	gcacccagag	tacaacagac	ccttgrtcgc	240
taacgacctc	atgctcatca	agttggacga	atrcgtgtcr	gagtcctgaca	ccatccggag	300
catcagcatt	gcttcgcagt	gccctaccgc	ggggaactct	tgcttcgltt	ctgcttgggg	360
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tgaggaggtc	tgagtaagr	tctatgaccc	gctgtaccac	cccagcatgt	tcttgcgccg	480
cggagggcaa	gaccagaagg	actccctgcaa	cgggtgactct	ggggggcccc	tgatctgcaa	540
cgggtacttg	cagggcccttg	tgtctttcgg	aaaagccccg	tgtggccaa	ttggcgtgcc	600
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agacccccca	gccccctcct	cctcagaccc	aggagtccag	ccctccctcc	ccaggagtc	900
ggagtcacag	ccccccagcc	cctccctcct	cagacccnagg	ggtcacaggcc	ccnaccct	960
cctccctcag	actcagaggt	ccaagcccc	aacccctcct	tcccagacc	cagaggtcca	1020
ggtcacagcc	cctccctcct	cagacccagc	ggtcacatgc	caactagact	ctccctgtac	1080
acagtgcctc	cttgtggcac	gttgacccaa	ccttaccagt	tggtttttca	tttttgtcc	1140
cttccctcta	gatccagaaa	tcaagttctaa	gagaagcgca	aaaaaanaa	aaaaaanaa	1200
aaaaaanaa	aaaaa					1215

&lt;210&gt; 327

&lt;211&gt; 220

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 327

Glu Asp Cys Ser Pro His Ser Gln Pro Trp Gln Ala Ala Leu Val Met	1	5	10	15
Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln Trp Val	20	25	30	35
Leu Ser Ala Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu Gly	35	40	45	50
Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val Glu	50	55	60	65
Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro Leu Leu Ala	65	70	75	80
Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu Ser Asp	85	90	95	100
Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala Gly Asn	100	105	110	115
Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Gly Arg Met Pro	115	120	125	130
Thr Val Leu Gln Cys Val Asn Val Ser Val Val Ser Glu Glu Val Cys	130	135	140	145
Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe Lys Ala Gly	145	150	155	160
Gly Gly Gln Asp Gln Lys Asp Ser Cys Asn Gly Asp Ser Gly Gly Pro	165	170	175	180
Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe Gly Lys Ala	180	185	190	195
Pro Cys Gly Gln Val Gly Val Pro Gly Val Tyr Thr Asn Leu Cys Lys	195	200	205	210
Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Ala Ser	210	215	220	

&lt;210&gt; 328



110

<211> 234  
 <212> DNA  
 <213> Homo sapien

<400> 328  
 cgtctgtctc tggtagctgc agccaaatca taaacggcga ggartgcagc ccgcactcgc 60  
 agcccttggca ggccgcaactg gtcattgaaa aggaattggt ctgctcgggc gtccctgggtgc 120  
 atccgcagtg ggtgctgtca gccacacact gtttccagaa ctctacacc atcgggctgg 180  
 gcctgcacag tcttgaggcc gaccaagagc cagggagcca gatggtggag gcc 234

<210> 329  
 <211> 77  
 <212> PRT  
 <213> Homo sapien

<400> 329  
 Leu Val Ser Gly Ser Cys Ser Gln Ile Ile Asn Gly Glu Asp Cys Ser  
 1 5 10 15  
 Pro His Ser Gln Pro Trp Gln Ala Ala Leu Val Met Glu Asn Glu Leu  
 20 25 30  
 Phe Cys Ser Gly Val Leu Val His Pro Gln Trp Val Leu Ser Ala Thr  
 35 40 45  
 His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu Gly Leu His Ser Leu  
 50 55 60  
 Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val Glu Ala  
 65 70 75

<210> 330  
 <211> 70  
 <212> DNA  
 <213> Homo sapien

<400> 330  
 cccaacacaa tggcccgatc ccattccctga ctccggccctc aggatcgctc gtctctggta 60  
 gctgcagcra 70

<210> 331  
 <211> 22  
 <212> PRT  
 <213> Homo sapien

<400> 331  
 Gln His Asn Gly Pro Ile Pro Ser Leu Thr Pro Pro Ser Gly Ser Leu  
 1 5 10 15  
 Val Ser Gly Ser Cys Ser  
 20

<210> 332  
 <211> 2507  
 <212> DNA  
 <213> Homo sapien

<400> 332  
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 tgccttctct tctgcatatg gctgcgcccc aaatcaggaa aatgctgtcc agtgggggtg 120

gtacatcaac	tgttcagctt	cctgggaag	tagttgtgt	cacaggagct	aatncaggt	180
tgggaagg	gacagccaa	gagctggctc	agagaggagc	tggagtatat	ttagcttgc	240
gggagctgga	aaaggggaa	ttggtaggca	aagagatcca	gaucacgaca	gggaaccagc	300
aggtgttgg	gcggaaactg	gacctgtctg	atactaagtc	tattcgagct	tttgctagg	360
gcttcttagc	tgaggaaag	caactccacg	ttttgatcaa	caatgcagg	gtgatgatgt	420
gtccgtactc	gaagaacga	gatggctttg	agatgcacat	aggagtcaac	caacttgggtc	480
acttctctct	ancccatctg	ctgctagaga	aactaaagga	atcagcccaa	tcaaggatag	540
taaatgtgtc	ttccctcgca	catcacttgg	gaaggatcca	cttccataac	ctgcaggggc	600
agaaatttca	caatgcaggc	ctggcctact	gtcacagcaa	gctagccaac	acccctcttca	660
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gcacagtcga	atrtgactg	gttcggcact	catctttcat	ggatggagtg	tgttggcttt	780
tctccctttt	catcaagact	cttcagcagg	gagccagac	cagcctgcac	tgtgccttaa	840
cagaaggtct	tgagattcta	agtgggaate	atttcagtga	ctgtcatgtg	gcatgggtct	900
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tyggctctcc	aatagactaa	caggcagtcg	cagttggacc	caagagagga	ctgcagcaga	1020
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agagagcaaa	accttccagc	cttgcctgct	tgtgttccag	ttaaaactca	gtgtactgoc	1140
agattcgtct	aaatgtctgt	catgtccaga	tctactttgc	ttctgttact	gccagagtta	1200
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&lt;210&gt; 333

&lt;211&gt; 3030

&lt;212&gt; DNA

&lt;213&gt; Homo Sapien

&lt;400&gt; 333

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&lt;210&gt; 334

&lt;211&gt; 2417

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 334

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&lt;210&gt; 335

&lt;211&gt; 2984

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 335

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&lt;210&gt; 336

&lt;211&gt; 147

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 336

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 Val Ile Glu Leu Glu Arg Lys Phe Ser His Gln Lys Tyr Leu Ser Ala  
 50 55 60  
 Pro Glu Arg Ala His Leu Ala Lys Asn Leu Lys Leu Thr Glu Thr Gln  
 65 70 75 80

Val Lys Ile Trp Phe Gln Asn Arg Arg Tyr Lys Thr Lys Arg Lys Gln  
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 Gly Ala Arg Val Tyr Leu Ala Cys Arg Asp Val Glu Lys Gly Glu Leu  
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 Arg Lys Leu Asp Leu Ser Asp Thr Lys Ser Ile Arg Ala Phe Ala Lys  
 100 105 110  
 Gly Phe Leu Ala Glu Glu Lys His Leu His Val Leu Ile Asn Asn Ala  
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 Ser Leu Ala His His Leu Gly Arg Ile His Phe His Asn Leu Gln Gly  
                                  180                      185                      190  
 Glu Lys Phe Tyr Asn Ala Gly Leu Ala Tyr Cys His Ser Lys Leu Ala  
                                  195                      200                      205  
 Asn Ile Leu Phe Thr Gln Glu Leu Ala Arg Arg Leu Lys Gly Ser Gly  
                                  210                      215                      220  
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 Arg His Ser Ser Phe Met Arg Trp Met Trp Trp Leu Phe Ser Phe Phe  
                                  245                                   250                                   255  
 Ile Lys Thr Pro Gln Gln Gly Ala Gln Thr Ser Leu His Cys Ala Leu  
                                  260                                   265                                   270  
 Thr Glu Gly Leu Glu Ile Leu Ser Gly Asn His Phe Ser Asp Cys His  
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 Val Ala Trp Val Ser Ala Gln Ala Arg Asn Glu Thr Ile Ala Arg Arg  
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 Leu Trp Asp Val Ser Cys Asp Leu Leu Gly Leu Pro Ile Asp  
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 <212> DNA  
 <213> Homo sapien

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 ctgatttcta acattgtctt taatgaccac aagacaacca acag 344

<210> 342  
 <211> 592  
 <212> DNA  
 <213> Homo sapien

&lt;400&gt; 342

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cctggcaggt	aaaccaatgc	caagagagtg	atggaaacca	tggcaagac	tttgttgatg	180
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aagtgccact	gtggaaagag	ttcctgtgtg	tgctgaagtt	ctgaaggcca	gtcaaattca	360
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cccggtgctt	tatgcacaata	atcgtcttct	tctaaatttc	tcttaggett	cattttccaa	480
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&lt;210&gt; 343

&lt;211&gt; 382

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 343

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ggggtagtgt	gaagggaactg	aaattgtggg	gggaaggctg	gaggacacac	antaagagg	360
aaaccaccaa	gctgaaaaaa	aa				382

&lt;210&gt; 344

&lt;211&gt; 536

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 344

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&lt;210&gt; 345

&lt;211&gt; 251

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 345

accttttgag	gtctctctca	ccacctccac	agccaccgtc	accgtgggat	gtgctggatg	60
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gtgccatttc	c					251



<210> 346  
 <211> 282  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (282)  
 <223> n = A, T, C or G

<400> 346  
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 agggagacta tacctggcgc ttgcctaaag tgagagggtct tccctccgc accaaaaaat 180  
 agaaaggctt tctatttcac tggcccaggc aggggggaagg agagtactt tgagctctg 240  
 ggtctcatt cccaaggcgc ctcaatgct catnaaaacc aa 282

<210> 347  
 <211> 201  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (1) ... (201)  
 <223> n = A, T, C or G

<400> 347  
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 taatataac tcttaanaa ttactanag cttttaccta agctcctaaa tgcttgtaaa 120  
 tctgagactg actggaccos cccagaccca gggcaaaagat acatgttacc atatcatctt 180  
 tataaagaat tctttttgt c 201

<210> 348  
 <211> 251  
 <212> DNA  
 <213> Homo sapien

<400> 348  
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 aggagacact cccagcatgg aggggggtt atcttttcat cctagggtcag gtctacaatg 180  
 ggggaaggte ttattataga actcccaada gcccaactca ctctgcccac ccaccgatg 240  
 gccctgectc c 251

<210> 349  
 <211> 251  
 <212> DNA  
 <213> Homo sapien

<400> 349  
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 cagaagggtc tgaactctac gtgtaccag agaactaat gcaattcatg catccactt 180  
 agcaattttg taanaatcaa gaaacagacc ccaagagtct ttcaagatga ggaanaatcca 240

actcctggtt t

251

&lt;210&gt; 350

&lt;211&gt; 908

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 350

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cggtcggaat	tgcctcgggt	atgatgacag	agaaaatgat	ctcttccctct	gtgacaccaa	180
cacctgtaaa	tttgatgggg	aatgtttaag	aattggagac	actgtgactt	gcgtctgtca	240
gttcaagtgc	aacaatgact	atgtgctgt	gtgtggctcc	aattggggaga	gctaccagaa	300
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ccacatacct	tgtccggaa	attacaatcg	cttctgcacg	catgggaagt	gtgagcattc	780
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aatgcag						908

&lt;210&gt; 351

&lt;211&gt; 472

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 351

ccagttattt	gcaagtggta	agagcctatt	taccataaat	aatactaaga	acraactraa	60
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&lt;210&gt; 352

&lt;211&gt; 251

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 352

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atacatggaa	aggaggggga	agccaaccca	gaaatgggct	ttctctaate	ctgggatacc	240
aataagrcac	a					251

&lt;210&gt; 353

&lt;211&gt; 436

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 353

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cacatttatgg tattttttact atactgatta tattttatcat gtgactttota atttataaat	120
gtatccaaaa gcaaaacagc agatatatcaa aattaaagag acagaaagata garatttaca	180
gataaggcca ttltatcatt gacaaatccaa atccaatara tttaaacatt tgggaatatg	240
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tcatgtctga raaggctctc ccttcaatgg ggatgcacaa ctccaaatgc cacacaaatg	360
ttaacagaat actagattca cactggaaag gggttaaga agaatattt ttctataaaa	420
gggtctctaa tgtagt	436

&lt;210&gt; 354

&lt;211&gt; 854

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 354

ccttttctag ttaccagth ttctgcagg atgctgggta gggagtgtct gcaggaggag	60
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atcagggaacc accctttggg tctatatatt gcttattctg catcttttga gtaagatcat	180
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ttaatgtcac acctacaggc actgggtctc tgccttcang tattttgltcc tcacttttagg	360
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caatgtaccc attttccctt ccaaatgtg aggggggggc ccgtgtgtt caaggctgtc	840
acacgggagt tcaag	854

&lt;210&gt; 355

&lt;211&gt; 676

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 355

gaaattaagt atgagctaaa ttccctgtta aaacttctag gggtagacaga tctcttcaac	60
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ctgttcttca taaggcacac tcataccaac acgatcttat tctgtggcaa gcttgcctct	300
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gcttaagaa aaccag	676

&lt;210&gt; 356

&lt;211&gt; 574

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 356

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&lt;210&gt; 357

&lt;211&gt; 393

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 357

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&lt;210&gt; 358

&lt;211&gt; 630

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 358

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caagccagag	gttctccac	aaacaaccag				630

&lt;210&gt; 359

&lt;211&gt; 620

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 359

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ctgtaaagat gtgacagtgt		620

&lt;210&gt; 360

&lt;211&gt; 431

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 360

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agattcttag t		430

&lt;210&gt; 361

&lt;211&gt; 351

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 361

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&lt;210&gt; 362

&lt;211&gt; 463

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 362

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&lt;210&gt; 363

&lt;211&gt; 653

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(653)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 363

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&lt;210&gt; 364

&lt;211&gt; 401

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 364

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&lt;210&gt; 365

&lt;211&gt; 356

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 365

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&lt;210&gt; 366

&lt;211&gt; 1851

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 366

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&lt;210&gt; 367

&lt;211&gt; 668

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 367

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&lt;210&gt; 368

&lt;211&gt; 1512

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 368

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&lt;210&gt; 369

&lt;211&gt; 1853

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 369

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&lt;210&gt; 370

&lt;211&gt; 2184

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 370

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<210> 371  
 <211> 1855  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (2)...(1855)  
 <223> n - A,T,C or G

<400> 371

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<210> 372  
 <211> 1059  
 <212> DNA  
 <213> Homo sapien

<400> 372

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<210> 373  
 <211> 1155  
 <212> DNA  
 <213> Homo sapien

<400> 373						
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ggcgtctctg	gagaccacga	cyactctgct	atgaagacac	tcaggaacaa	gatgggcaag	300
tgggtgctgcc	actgcttccc	ctgctgcagg	gggagcggca	agagcaaggt	ggcgccttgg	360
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gocagagagt	atgctgtttc	tagtcatcat	catgtaattt	gccagttact	ttctgactac	1080
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<210> 374  
 <211> 2000  
 <212> DNA  
 <213> Homo sapien

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ggcgtctctg	gagaccacga	cyactctgct	atgaagacac	tcaggaacaa	gatgggcaag	300
tgggtgctgcc	actgcttccc	ctgctgcagg	gggagcggca	agagcaaggt	ggcgccttgg	360
ggagactatg	atgacagtgc	cttcatggag	ccuaggtacc	actgctggtg	agaagatctg	420
gacaagctcc	acagagctgc	ctgggtgggt	aaagtcccc	gaaaggtatc	catcgtcatg	480
ctcagggaca	ctgacgtgaa	caugaaggac	aaagcaaaaga	ggactgctct	acatctggcc	540

tctgccaatg	ggaatccaga	agttagtaaaa	ctcctgctgg	acagacgatg	tcaacttaant	600
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tgtgctgtaa	tgttgctgga	acatggcact	gatccaaata	ttccagatga	gtatggaaat	720
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gccagagagt	atgctgcttc	tagtcctcat	catgtaattt	gccagttact	ttctgactac	1080
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&lt;210&gt; 375

&lt;211&gt; 2040

&lt;212&gt; DNA

&lt;213&gt; Homo sapien

&lt;400&gt; 375

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gtcagccttc	tacttgagca	aaatattgat	gtatcttctc	aagatctatc	tggacagacg	1020
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gaaatgaaga agcacggaag tactcatgtc ggattccag aaaacctgac taatgggtgcr      1680
actgctggca atgggtgatga tggattaatt cctccaagga agagcagaac arctgaaagc      1740
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aagcaatttt ytgangaaca gaacactgga atattacag atgagattct gattcatgaa      1860
gaaaagcaga tagaagtggg tgaaaaaatg nattctgagc ttctctcttag ttgtaagaaa      1920
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&lt;210&gt; 376

&lt;211&gt; 329

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 376

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Leu His Leu Ala Gly Ser Asp Leu Leu Ser Arg Ser Leu Met Ala Glu
 20          25          30
Glu Tyr Thr Ile Val His Ala Ser Phe Ile Ser Cys Ile Ser Ser Ser
 35          40          45
Leu Asp Gly Gln Gly Glu Arg Gln Glu Arg Gly His Phe Trp Arg
 50          55          60
Pro Gln Arg Leu Leu Cys Glu Asp Ala Trp Glu Gln Glu Val Gln Val
 65          70          75          80
Val Leu Pro Leu Leu Pro Leu Leu Gln Gly Ser Gly Lys Ser Asn Val
 85          90          95
Val Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe Met Asp Pro Arg Tyr
100          105          110
His Val His Gly Glu Asp Leu Asp Lys Leu His Arg Ala Ala Trp Trp
115          120          125
Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met Leu Arg Asp Thr Asp
130          135          140
Val Asn Lys Arg Asp Lys Gln Lys Arg Thr Ala Leu His Leu Ala Ser
145          150          155          160
Ala Asn Gly Asn Ser Glu Val Val Lys Leu Val Leu Asp Arg Arg Cys
165          170          175
Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr Ala Leu Thr Lys Ala
180          185          190
Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met Leu Leu Glu His Gly
195          200          205
Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn Thr Thr Leu His Tyr
210          215          220
Ala Val Tyr Asn Glu Asp Lys Leu Met Ala Lys Ala Leu Leu Leu Tyr
225          230          235          240
Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly Leu Thr Pro Leu Leu
245          250          255
Leu Gly Ile His Glu Gln Lys Gln Gln Val Val Lys Phe Leu Ile Lys
260          265          270
Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr Gly Arg Thr Ala Leu
275          280          285
Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile Val Ser Pro Leu Leu

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290                      295                      300  
 Glu Gln Asn Val Asp Val Ser Ser Gln Asp Leu Glu Arg Arg Pro Glu  
 305                      310                      315                      320  
 Ser Met Leu Phe Leu Val Ile Ile Met  
                          325

<210> 377  
 <211> 148  
 <212> PRT  
 <213> Homo sapien

<220>  
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                          20                      25                      30  
 Asp Leu Ile Val Met Leu Arg Asp Thr Asp Val Asn Lys Xaa Asp Lys  
                          35                      40                      45  
 Gln Lys Arg Thr Ala Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu  
                          50                      55                      60  
 Val Val Lys Leu Xaa Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp  
 65                      70                      75                      80  
 Asn Lys Lys Arg Thr Ala Leu Xaa Lys Ala Val Gln Cys Gln Glu Asp  
                          85                      90                      95  
 Glu Cys Ala Leu Met Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro  
                          100                      105                      110  
 Asp Glu Tyr Gly Asn Thr Thr Leu His Tyr Ala Xaa Tyr Asn Glu Asp  
                          115                      120                      125  
 Lys Leu Met Ala Lys Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser  
                          130                      135                      140  
 Lys Asn Lys Val  
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<210> 378  
 <211> 1719  
 <212> PRT  
 <213> Homo sapien

<400> 378  
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                          20                      25                      30  
 Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp  
                          35                      40                      45  
 His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Trp  
                          50                      55                      60  
 Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val  
 65                      70                      75                      80  
 Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn

	85		90		95
Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys Cys Arg Gly Ser					
	100		105		110
Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe					
	115		120		125
Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His					
	130		135		140
Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met					
	145		150		155
Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln Lys Arg Thr Ala					
	165		170		175
Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu					
	180		185		190
Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr					
	195		200		205
Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met					
	210		215		220
Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn					
	225		230		235
Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys					
	245		250		255
Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly					
	260		265		270
Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys Gln Gln Val Val					
	275		280		285
Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr					
	290		295		300
Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile					
	305		310		315
Val Ser Leu Leu Leu Glu Gln Asn Ile Asp Val Ser Ser Gln Asp Leu					
	325		330		335
Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val					
	340		345		350
Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln Met Leu Lys Ile					
	355		360		365
Ser Ser Glu Asn Ser Asn Pro Glu Asn Val Ser Arg Thr Arg Asn Lys					
	370		375		380
Pro Arg Thr His Met Val Val Glu Val Asp Ser Met Pro Ala Ala Ser					
	385		390		395
Ser Val Lys Lys Pro Phe Gly Leu Arg Ser Lys Met Gly Lys Trp Cys					
	405		410		415
Cys Arg Cys Phe Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly					
	420		425		430
Thr Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys					
	435		440		445
Met Gly Lys Trp Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly					
	450		455		460
Lys Ser Asn Val Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys					
	465		470		475
Thr Leu Arg Asn Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys					
	485		490		495
Cys Arg Gly Ser Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp					
	500		505		510
Asp S r Ala Phe Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu					
	515		520		525

Asp Lys Leu His Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp  
 530 535 540  
 Leu Ile Val Met Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln  
 545 550 555 560  
 Lys Arg Thr Ala Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val  
 565 570 575  
 Val Lys Leu Leu Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn  
 580 585 590  
 Lys Lys Arg Thr Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu  
 595 600 605  
 Cys Ala Leu Met Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp  
 610 615 620  
 Glu Tyr Gly Asn Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys  
 625 630 635 640  
 Leu Met Ala Lys Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys  
 645 650 655  
 Asn Lys His Gly Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys  
 660 665 670  
 Gln Gln Val Val Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala  
 675 680 685  
 Leu Asp Arg Tyr Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly  
 690 695 700  
 Ser Ala Ser Ile Val Ser Leu Leu Leu Glu Gln Asn Ile Asp Val Ser  
 705 710 715 720  
 Ser Gln Asp Leu Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser  
 725 730 735  
 His His His Val Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln  
 740 745 750  
 Met Leu Lys Ile Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys  
 755 760 765  
 Leu Thr Ser Glu Glu Glu Ser Gln Arg Phe Lys Gly Ser Glu Asn Ser  
 770 775 780  
 Gln Pro Glu Lys Met Ser Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp  
 785 790 795 800  
 Arg Glu Val Glu Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly  
 805 810 815  
 Leu Leu Glu Asn Leu Thr Asn Gly Val Thr Ala Gly Asn Gly Asp Asn  
 820 825 830  
 Gly Leu Ile Pro Gln Arg Lys Ser Arg Thr Pro Glu Asn Gln Gln Phe  
 835 840 845  
 Pro Asp Asn Glu Ser Glu Glu Tyr His Arg Ile Cys Glu Leu Val Ser  
 850 855 860  
 Asp Tyr Lys Glu Lys Gln Met Pro Lys Tyr Ser Ser Glu Asn Ser Asn  
 865 870 875 880  
 Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Leu  
 885 890 895  
 Glu Gly Ser Glu Asn Gly Gln Pro Glu Leu Glu Asn Phe Met Ala Ile  
 900 905 910  
 Glu Glu Met Lys Lys His Gly Ser Thr His Val Gly Phe Pro Glu Asn  
 915 920 925  
 Leu Thr Asn Gly Ala Thr Ala Gly Asn Gly Asp Asp Gly Leu Ile Pro  
 930 935 940  
 Pro Arg Lys Ser Arg Thr Pro Glu Ser Gln Gln Phe Pro Asp Thr Glu  
 945 950 955 960  
 Asn Glu Glu Tyr His Ser Asp Glu Gln Asn Asp Thr Gln Lys Gln Phe



965 970 975  
 Cys Glu Glu Gln Asn Thr Gly Ile Leu His Asp Glu Ile Leu Ile His  
 980 985 990  
 Glu Glu Lys Gln Ile Glu Val Val Glu Lys Met Asn Ser Glu Ile Ser  
 995 1000 1005  
 Leu Ser Cys Lys Lys Glu Lys Asp Ile Leu His Glu Asn Ser Thr Leu  
 1010 1015 1020  
 Arg Glu Glu Ile Ala Met Leu Arg Leu Glu Ile Asp Thr Met Lys His  
 1025 1030 1035 104  
 Gln Ser Gln Leu Pro Arg Thr His Met Val Val Glu Val Asp Ser Met  
 1045 1050 1055  
 Pro Ala Ala Ser Ser Val Lys Lys Pro Phe Gly Leu Arg Ser Lys Met  
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 Gly Lys Trp Cys Cys Arg Cys Phe Pro Cys Cys Arg Glu Ser Gly Lys  
 1075 1080 1085  
 Ser Asn Val Gly Thr Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr  
 1090 1095 1100  
 Leu Arg Ser Lys Met Gly Lys Trp Cys Arg His Cys Phe Pro Cys Cys  
 1105 1110 1115 112  
 Arg Gly Ser Gly Lys Ser Asn Val Gly Ala Ser Gly Asp His Asp Asp  
 1125 1130 1135  
 Ser Ala Met Lys Thr Leu Arg Asn Lys Met Gly Lys Trp Cys Cys His  
 1140 1145 1150  
 Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Lys Val Gly Ala Trp  
 1155 1160 1165  
 Gly Asp Tyr Asp Asp Ser Ala Phe Met Glu Pro Arg Tyr His Val Arg  
 1170 1175 1180  
 Gly Glu Asp Leu Asp Lys Leu His Arg Ala Ala Trp Trp Gly Lys Val  
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 Pro Arg Lys Asp Leu Ile Val Met Leu Arg Asp Thr Asp Val Asn Lys  
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 Asn Ser Glu Val Val Lys Leu Leu Leu Asp Arg Arg Cys Gln Leu Asn  
 1235 1240 1245  
 Val Leu Asp Asn Lys Lys Arg Thr Ala Leu Ile Lys Ala Val Gln Cys  
 1250 1255 1260  
 Gln Glu Asp Glu Cys Ala Leu Met Leu Leu Glu His Gly Thr Asp Pro  
 1265 1270 1275 128  
 Asn Ile Pro Asp Glu Tyr Gly Asn Thr Thr Leu His Tyr Ala Ile Tyr  
 1285 1290 1295  
 Asn Glu Asp Lys Leu Met Ala Lys Ala Leu Leu Leu Tyr Gly Ala Asp  
 1300 1305 1310  
 Ile Glu Ser Lys Asn Lys His Gly Leu Thr Pro Leu Leu Gly Val  
 1315 1320 1325  
 His Glu Gln Lys Gln Gln Val Val Lys Phe Leu Ile Lys Lys Lys Ala  
 1330 1335 1340  
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 Val Cys Cys Gly Ser Ala Ser Ile Val Ser Leu Leu Leu Glu Gln Asn  
 1365 1370 1375  
 Ile Asp Val Ser Ser Gln Asp Leu Ser Gly Gln Thr Ala Arg Glu Tyr  
 1380 1385 1390  
 Ala Val Ser Ser His His His Val Ile Cys Gln Leu Leu Ser Asp Tyr  
 1395 1400 1405

Lys Glu Lys Gln Met Leu Lys Ile Ser Ser Glu Asn Ser Asn Pro Glu  
 1410 1415 1420  
 Gln Asp Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Phe Lys Gly  
 1425 1430 1435 144  
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 1445 1450 1455  
 Lys Asp Gly Asp Arg Glu Val Glu Glu Glu Met Lys Lys His Glu Ser  
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 1475 1480 1485  
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 1490 1495 1500  
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 1505 1510 1515 152  
 Glu Leu Val Ser Asp Tyr Lys Glu Lys Gln Met Pro Lys Tyr Ser Ser  
 1525 1530 1535  
 Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu Glu Glu  
 1540 1545 1550  
 Ser Gln Arg Leu Glu Gly Ser Glu Asn Gly Gln Pro Glu Lys Arg Ser  
 1555 1560 1565  
 Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp Arg Glu Leu Glu Asn Phe  
 1570 1575 1580  
 Met Ala Ile Glu Glu Met Lys Lys His Gly Ser Thr His Val Gly Phe  
 1585 1590 1595 160  
 Pro Glu Asn Leu Thr Asn Gly Ala Thr Ala Gly Asn Gly Asp Asp Gly  
 1605 1610 1615  
 Leu Ile Pro Pro Arg Lys Ser Arg Thr Pro Glu Ser Gln Gln Phe Pro  
 1620 1625 1630  
 Asp Thr Glu Asn Glu Glu Tyr His Ser Asp Glu Gln Asn Asp Thr Gln  
 1635 1640 1645  
 Lys Gln Phe Cys Glu Glu Gln Asn Thr Gly Ile Leu His Asp Glu Ile  
 1650 1655 1660  
 Leu Ile His Glu Glu Lys Gln Ile Glu Val Val Glu Lys Met Asn Ser  
 1665 1670 1675 168  
 Glu Leu Ser Leu Ser Cys Lys Lys Glu Lys Asp Ile Leu His Glu Asn  
 1685 1690 1695  
 Ser Thr Leu Arg Glu Glu Ile Ala Met Leu Arg Leu Glu Leu Asp Thr  
 1700 1705 1710  
 Met Lys His Gln Ser Gln Leu  
 1715

<210> 379  
 <211> 656  
 <212> PRT  
 <213> Homo sapien

<400> 379

Met Val Val Glu Val Asp Ser Met Pro Ala Ala Ser Ser Val Lys Lys  
 1 5 10 15  
 Pro Phe Gly Leu Arg Ser Lys Met Gly Lys Trp Cys Cys Arg Cys Phe  
 20 25 30  
 Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp  
 35 40 45  
 His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Trp  
 50 55 60

Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val  
 65 70 75 80  
 Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn  
 85 90 95  
 Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys Cys Arg Gly Ser  
 100 105 110  
 Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe  
 115 120 125  
 Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His  
 130 135 140  
 Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met  
 145 150 155 160  
 Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln Lys Arg Thr Ala  
 165 170 175  
 Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu  
 180 185 190  
 Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr  
 195 200 205  
 Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met  
 210 215 220  
 Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn  
 225 230 235 240  
 Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys  
 245 250 255  
 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly  
 260 265 270  
 Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys Gln Gln Val Val  
 275 280 285  
 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr  
 290 295 300  
 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile  
 305 310 315 320  
 Val Ser Leu Leu Leu Glu Gln Asn Ile Asp Val Ser Ser Gln Asp Leu  
 325 330 335  
 Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val  
 340 345 350  
 Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln Met Leu Lys Ile  
 355 360 365  
 Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu  
 370 375 380  
 Glu Glu Ser Gln Arg Phe Lys Gly Ser Glu Asn Ser Gln Pro Glu Lys  
 385 390 395 400  
 Met Ser Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp Arg Glu Val Glu  
 405 410 415  
 Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly Leu Leu Glu Asn  
 420 425 430  
 Leu Thr Asn Gly Val Thr Ala Gly Asn Gly Asp Asn Gly Leu Ile Pro  
 435 440 445  
 Gln Arg Lys Ser Arg Thr Pro Glu Asn Gln Gln Phe Pro Asp Asn Glu  
 450 455 460  
 Ser Glu Glu Tyr His Arg Ile Cys Glu Leu Val Ser Asp Tyr Lys Glu  
 465 470 475 480  
 Lys Gln Met Pro Lys Tyr Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp  
 485 490 495  
 Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Leu Glu Gly Ser Glu

500	505	510
Asn Gly Glu Pro Glu Leu Glu	Asn Phe Met Ala Ile Glu	Glu Glu Met Lys
515	520	525
Lys His Gly Ser Thr His Val Gly	Phe Pro Glu Asn Leu Thr Asn Gly	
530	535	540
Ala Thr Ala Gly Asn Gly Asp Asp	Gly Leu Ile Pro Pro Arg Lys Ser	
545	550	555
Arg Thr Pro Glu Ser Gln Gln Phe	Pro Asp Thr Glu Asn Glu Glu Tyr	
565	570	575
His Ser Asp Glu Gln Asn Asp Thr	Gln Lys Gln Phe Cys Glu Glu Gln	
580	585	590
Asn Thr Gly Ile Leu His Asp Glu	Ile Leu Ile His Glu Glu Lys Gln	
595	600	605
Ile Glu Val Val Glu Lys Met Asn	Ser Glu Leu Ser Leu Ser Cys Lys	
610	615	620
Lys Glu Lys Asp Ile Leu His Glu	Asn Ser Thr Leu Arg Glu Glu Ile	
625	630	635
Ala Met Leu Arg Leu Glu Leu Asp	Thr Met Lys His Gln Ser Gln Leu	
645	650	655

<210> 380  
 <211> 671  
 <212> PRT  
 <213> Homo sapien

<400> 380
Met Val Val Glu Val Asp Ser Met Pro Ala Ala Ser Ser Val Lys Lys
1 5 10 15
Pro Phe Gly Leu Arg Ser Lys Met Gly Lys Trp Cys Cys Arg Cys Phe
20 25 30
Pro Cys Cys Arg Glu Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp
35 40 45
His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Trp
50 55 60
Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val
65 70 75 80
Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn
85 90 95
Lys Met Gly Lys Trp Cys Cys His Cys Phe Pro Cys Cys Arg Gly Ser
100 105 110
Gly Lys Ser Lys Val Gly Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe
115 120 125
Met Glu Pro Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His
130 135 140
Arg Ala Ala Trp Trp Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met
145 150 155 160
Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Gln Lys Arg Thr Ala
165 170 175
Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu
180 185 190
Leu Asp Arg Arg Cys Gln Leu Asn Val Leu Asp Asn Lys Lys Arg Thr
195 200 205
Ala Leu Ile Lys Ala Val Gln Cys Gln Glu Asp Glu Cys Ala Leu Met
210 215 220
Leu Leu Glu His Gly Thr Asp Pro Asn Ile Pro Asp Glu Tyr Gly Asn

225 230 235 240  
 Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys  
 245 250 255  
 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly  
 260 265 270  
 Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys Gln Gln Val Val  
 275 280 285  
 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr  
 290 295 300  
 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile  
 305 310 315 320  
 Val Ser Leu Leu Leu Glu Gln Asn Ile Asp Val Ser Ser Gln Asp Leu  
 325 330 335  
 Ser Gly Gln Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val  
 340 345 350  
 Ile Cys Gln Leu Leu Ser Asp Tyr Lys Glu Lys Gln Met Leu Lys Ile  
 355 360 365  
 Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp Leu Lys Leu Thr Ser Glu  
 370 375 380  
 Glu Glu Ser Gln Arg Phe Lys Gly Ser Glu Asn Ser Gln Pro Glu Lys  
 385 390 395 400  
 Met Ser Gln Glu Pro Glu Ile Asn Lys Asp Gly Asp Arg Glu Val Glu  
 405 410 415  
 Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly Leu Leu Glu Asn  
 420 425 430  
 Leu Thr Asn Gly Val Thr Ala Gly Asn Gly Asp Asn Gly Leu Ile Pro  
 435 440 445  
 Glu Arg Lys Ser Arg Thr Pro Glu Asn Gln Gln Phe Pro Asp Asn Glu  
 450 455 460  
 Ser Glu Glu Tyr His Arg Ile Cys Glu Leu Val Ser Asp Tyr Lys Glu  
 465 470 475 480  
 Lys Gln Met Pro Lys Tyr Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp  
 485 490 495  
 Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Leu Glu Gly Ser Glu  
 500 505 510  
 Asn Gly Gln Pro Glu Lys Arg Ser Gln Glu Pro Glu Ile Asn Lys Asp  
 515 520 525  
 Gly Asp Arg Glu Leu Glu Asn Phe Met Ala Ile Glu Glu Met Lys Lys  
 530 535 540  
 His Gly Ser Thr His Val Gly Phe Pro Glu Asn Leu Thr Asn Gly Ala  
 545 550 555 560  
 Thr Ala Gly Asn Gly Asp Asp Gly Leu Ile Pro Pro Arg Lys Ser Arg  
 565 570 575  
 Thr Pro Glu Ser Gln Gln Phe Pro Asp Thr Glu Asn Glu Glu Tyr His  
 580 585 590  
 Ser Asp Glu Gln Asn Asp Thr Gln Lys Gln Phe Cys Glu Glu Gln Asn  
 595 600 605  
 Thr Gly Ile Leu His Asp Glu Ile Leu Ile His Glu Glu Lys Gln Ile  
 610 615 620  
 Glu Val Val Glu Lys Met Asn Ser Glu Leu Ser Leu Ser Cys Lys Lys  
 625 630 635 640  
 Glu Lys Asp Ile Leu His Glu Asn Ser Thr Leu Arg Glu Glu Ile Ala  
 645 650 655  
 Met Leu Arg Leu Glu Leu Asp Thr Met Lys His Gln Ser Gln Leu  
 660 665 670

<210> 381  
 <211> 251  
 <212> DNA  
 <213> Homo sapien

<400> 381

ggagagagcgt ctgctggggc aggaaggggt ttccctgccc tctcactgt cctcaccac	60
ggtaacatgc ttccctcag ggtatccca cccaggggc ttaccatgac ctclgagggg	120
cccattatccc agyagaagca ttggggaggt gggggcaggt gaaggaccca ggactcacac	180
atcctggggc tccaaggcag agggaggggt cctcaagaaq gt caggagga aaatccgtaa	240
caagcagta g	251

<210> 382  
 <211> 3279  
 <212> DNA  
 <213> Homo sapiens

<400> 382

cttccctgcag ccccccctgct ggtgaggggc ccggggcagga acagtggacc caacatggaa	60
ctgctgaggg gtgtcaggaa gtgtcgggc tctggggcag gggggagggg tggggagtgt	120
cactgggagg ggacatcctg cagaaggtag gaghgagcaa acacccgctg caggggaggg	180
gagagccctg cggcacctgg gggagcagag gtagcagrac ctgcccagga ctgggaggaq	240
gggctcggag ggcgtgagga gtagcaggg ggtgcctgg ctggagtgg ggcacaggg	300
cagggcgcca gattggctca cacagggaag agagggccc tctgcaggg cctcactgg	360
gcccacaggag gacactgctt ttctctgag gagt caggag ctgtggatgg tgcaggacag	420
aagaaggana gggcctggct caggtgtcca gaggctgtcg ctggcttccc ttggggatca	480
gactcaggg agggagggcg gcagggtgt gggggagtg ccagctgagga tgactgggg	540
gtggctccag gcttgcctcc tgcctgggc ctccccagc cccctcaca gctcctggc	600
cctcagttct tccctcacc tccctcctcc atctggctc agtgggtcat tctgatcact	660
gaactgacca taccacagcc tgcctcaggg cctccatggc tccccaatgc cctggagagg	720
ggacatctag tccagagagta gtcctgagga ggtggcctct gcatgtgacc tgtggggcga	780
gcactctgca gatggctccc gcccctcacc tgcctgacct totgacggga ctgtcctcct	840
ggacattgac cttctgtgag gaggcggcc ctgaagtcct cttcccatag gccagactg	900
gagccttgtt cctctgttly gactcctgc ccatattctt gtgggagtg gtcttggaga	960
catctctgtc tctcctgag agctgggaat tgcctcagt catctgctg ccgggttctg	1020
agagatggag ttgctaggc agttattgg gccaatctt tccactgtgt ctctcctcct	1080
ttacctcag ggtgattctg ggggtccact tctctgtaat ggtgtgcttc aaggtatcac	1140
atcatggggc cctgagccat gtcacctgcc tgaagaagct gctgtgtaca ccaaggtgtt	1200
gcattaccgg aagtggatca aggacacat cgcagccaac cctggagtc cctgtccc	1260
ccctacctc tagtaaatlt aagtccacct cacttctgg catcatttg ccttcttgg	1320
tgttgagac ctgaggttg gaactcact ggcgaagct ctagcctcct gactcctact	1380
gactgtgtt ttctggtgt ggtccagg ctgctaggaa aaggaatgg cagacacagg	1440
tgtatgccaa tgtttctgaa elgggtataa ttctgctct ccttuggaa cactggctgt	1500
ctctgagag ttctcgtca gtttcagtga ggaacacac aaagacgtgg gtgaccatgt	1560
tgtttgtgg gtgcagagat gggaggggtg gggccacccc tggagagtg gacagtga	1620
caaggtggac actctctaca gatcactgag gataagctgg agccacatg catgaggcac	1680
acacacagca aggttgaguc tgtaaacata gccacagctg tctgggggc actgggaagc	1740
ctagataagg ccgtgagcag aaagaaaggg aggtatctcc tatgttgtt aaggagggac	1800
tagggggaga aactgaaagc tgattaatc caggaggtt gtccaggtcc cccaaaccac	1860
cgtcagatt gatgatttcc tagcaggact tacagaanta aagagctate atgctgtgtt	1920
ttattatgg ttgttccatt gataggalac atactgaant cagcaaccaa aacagatgt	1980
tagattagag tgtggagaa acagagggaa acttgcaglt acgagactg gcaacttggc	2040
ttactaagt ttccagactg gcagggaagt aaacctatta ggtgaggaac cttgtggagt	2100
gtagctgac cagctgalt agyaaactagc cagggtgggg ccttccctt tggatggagg	2160

```

gcataatcga cagttattct ctccaaagtgg agacttaagg acagcatata attctccctg 2220
caaggatgta tgaataatag tacaaggtaa ttccaaactga ggaagctcac ctgatacetta 2280
gtgtccagggt tttttactgg gggctctgtag gacgagtatg ggaatactga ataattgacc 2340
tgaagtcttc agacctgaggt ttcccttagag ttccaaacaga tacagcatgg tccagagctcc 2400
cagatgtaca aaaacaggga ttcatcacaa atcccatctt tagaatgaag ggtctggcat 2460
ggcccaaggc cccaaagtata tcaaggcact tgggcagacc atgccaaggga atcaaatgtc 2520
atctcccagg agttattcaa gggctgagccc ttactctggg atgtacaggc tttagagcgt 2580
gcagggtctgc tgaagtcaccc ttttattgta cagaggatga gggaaaggga gaggatgagg 2640
aagccccctt ggggatttgg tttgggtctg tgatcagggt gtctatgggg ctatccctac 2700
aagaagagac ccagaaatag gggcacattg aggaatgata ctgagcccaa agagcattca 2760
atcattgttt tatttgcctt ctcttcacac cattgggtgag ggaaggatta ccacctggy 2820
gttatgaaga tggttgaaac cccacacat agcacctggg atatgagatc aacagtttct 2880
tagccataga gattcacagc ccagagcagg aggacgctgc acacatgca ggaagacatg 2940
ggggatgcgc tgggatttgg tgtgaagaag cagagactgt tagaggcagg ctttatagta 3000
acaagacggt ggggcaact ctggttccg tgggggaatg tcatggtctt gctttactna 3060
gttttgagac tggcaggtag tgaactcat taggctgaga accttgtgga atgacgtga 3120
ccagctgat agaggaagta gccaggtggg agccttccg agtgggtgtg ggacatctct 3180
ggcaagattt tgtggcactc ctggttacag atactggggc agcaaataaa ctgcaatctt 3240
gttttcagac cttaaaaaa aaaaaaaa aaaggtttt 3279

```

&lt;210&gt; 383

&lt;211&gt; 155

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 383

```

Met Ala Gly Val Arg Asp Gln Gly Gln Gly Ala Arg Trp Pro His Thr
      5                                10                    15

```

```

Gly Lys Arg Gly Pro Leu Leu Gln Gly Leu Thr Trp Ala Thr Gly Gly
      20                                25                    30

```

```

His Cys Phe Ser Ser Glu Gln Ser Gly Ala Val Asp Gly Ala Gly Gln
      35                                40                    45

```

```

Lys Lys Asp Arg Ala Trp Leu Arg Cys Pro Glu Ala Val Ala Gly Phe
      50                                55                    60

```

```

Pro Leu Gly Ser Asp Cys Arg Glu Gly Gly Arg Gln Gly Cys Gly Gly
      65                                70                    75                    80

```

```

Ser Asp Asp Glu Asp Asp Leu Gly Val Ala Pro Gly Leu Ala Pro Ala
      85                                90                    95

```

```

Trp Ala Leu Thr Gln Pro Pro Ser Gln Ser Pro Gly Pro Gln Ser Leu
     100                                105                    110

```

```

Pro Ser Thr Pro Ser Ser Ile Trp Pro Gln Trp Val Ile Leu Ile Thr
     115                                120                    125

```

```

Glu Leu Thr Ile Pro Ser Pro Ala His Gly Pro Pro Trp Leu Pro Asn
     130                                135                    140

```

```

Ala Leu Glu Arg Gly His Leu Val Arg Glu
     145                                150

```

<210> 384  
 <211> 557  
 <212> DNA  
 <213> Homo sapiens

<400> 384  
 ggatcctcta gagcggcgcg ctactactac taaattcgcg gercgcgcga cgaagaagag 60  
 aaagatgtgt ttgtgtttgg autctctgtg gtercttcca atgctgtggg ttcccaacca 120  
 ggggaagggt cccitttgca ttgccaagtg ccataccat gacactact ctaccatgg 180  
 tctgcctcct ggccaagcag gctggtttgc aagaatgaa tgaatgatc tacagctagg 240  
 acttaacctt gaantggaaa gtcttgcact cccatttgca ggatccgct gtgcacatgc 300  
 ctctgtagag agcagcattc cuagggaact tggaaacagt tggcactgta aggtgcttgc 360  
 tccccaaggc acatcctaaa aggtgttgtt atggtgaaa cgtcttccct ctttattgcc 420  
 ccttcttatt tatgtgaaca actgtttgtc ttttttgtt tttttttta actgttaagt 480  
 tcaattgtga aatggaatat catgcaata aattatgoga ttttttttc aaagtaaaa 540  
 aaaaaaaa 557

<210> 385  
 <211> 337  
 <212> DNA  
 <213> Homo sapiens

<400> 385  
 ttcccaggat atgtgcgagg gaagacacct tcaatctct tgaaggggct gattccttta 60  
 gtttctctag cagcagatgg gctaggagga agtgaccca gtaggttgat ctatgtgca 120  
 tctcaagacc atctgtgtgc ttcgagtacg gacacatcat cactcctgca ttgttgatca 180  
 aaacgtggag gtgttttccc tcagctaaag agcccttagc aaaaagtoga atagacttag 240  
 tatcagacag gtccagtttc cgcaccaaca cctgctggtt cctgtcgtg gcttggatct 300  
 gtttggccac caattccccc tttcccaat ccggga 337

<210> 386  
 <211> 300  
 <212> DNA  
 <213> Homo sapiens

<400> 386  
 gggcccgcta ccggccragg ccccgccctg cgagtcctcc tccrccgggt cctgcccgc 60  
 gcccgctcgg cccagagggt gggcggggg ctgctctac cggctggcgg ctgttaactca 120  
 gcgacotttg cccgaaggct ctaggcagga cccatcgacc ccagccgagg cggcggggc 180  
 gcggaatttg cccggtgtgt gggcgggagc ggaatgcgtg tccgaggag ggcagcgaag 240  
 atgttagcct tcgtgncag gaccgtggac agatccagg gctgtggtgt aacctcagcc 300

<210> 387  
 <211> 537  
 <212> DNA  
 <213> Homo sapiens

<400> 387  
 gggccgaagt gggcaccag ggactcttga caggcttctc tctcgggato atcaaggctg 60  
 cccctctctg tggcatcat atcagcact atgagttcgg caaaagcttc ttccagaggc 120  
 tgaaccaaag ccggttctg ggcggctgaa aggggcaagg aggcaggac cccgtctctc 180  
 ccacggatgg gggaggggac gaggagacc cagccaagtg ctttttctc agcautgagg 240  
 gagggggctt gtttccctt cctccggggc aaagctcca gggcagggcl gtcctctctg 300



```

ggggcccagc atttctctag acacaacttc ttctctgtgc tccagtcgtg gggatcaten 360
cttaccacacn ccccaagttc aagacccaat ctccagctg cccctctcgt gtttccctgt 420
gtttgtctga gctgggcatg tctccaggaa ccaaggaagc ctacgcctgg tctagtcctc 480
ctgacccctg ttaattccct aagtcctaaag atgatgaact tcaaaaaaaa aaaaaaa 537

```

&lt;210&gt; 388

&lt;211&gt; 520

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 388

```

aggataattt ttaaaccaat caaatgaaa aaacaacaa acaaaaaagg aattgtcatg 60
tgagggttaa ccagtttgca tccccctaat gtgggaaaag taagaggact actcaggaact 120
gtttgaagat tgcctctctt acagcttctg agaatttgtt tatttccctt gcccaagtga 180
ggaccccttc ccacacatgc cccagccca cctlaagcat ggtcccttgc caccaggaac 240
ccaggaaact gctacttgtg gactctacca gagaccagga ggggttggtt agctcacagg 300
acttccccc cccagaaga ttagcatccr atactgaact cactactaac tcaactagga 360
tcatactcaa ttgatggta tttagcaatt ccatttcttt ctgggttact taacacagaa 420
atcttctctc ttctcattac cagtaaaagg tcttggtatc tttctgttgg aatgatttct 480
atgaacttgt cttattttta tgggtgggtt tttttctggt 520

```

&lt;210&gt; 389

&lt;211&gt; 365

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 389

```

cgttgcctca gtttgacaga aggaagggcg gagcttcttc aaagtctaga gggagtgagg 60
gagtttaaggc tggatttcag atctgcctcg ttccagccgc agtgtgacct ctgtccccc 120
aargacttcc caataatct cccagcgcc tccagctca ggcgtctag aagcgtcttg 180
aagcctatgg ccagctgtct ttgtgtccc tctcaccgc ctgtctcar agctgaagct 240
cccaggaaac ctccagacta ccttctcttg ccttcagca ggggcttgc ccacattctc 300
tgagggtcag tggagaacc tagactccca ttgctagagg tagaaagggg aagggtgctg 360
gggag 365

```

&lt;210&gt; 390

&lt;211&gt; 221

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; [1]... (221)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 390

```

tgctctcca tcttgccccc gaattctctg tcaggaaagt ggggattggc cccatctgca 60
tacacggmtt ctcatgggtg tggacatct ctgtctggcg ttccaggaa gctctggct 120
gctctangag tctganenga nctgttgc cactntgaca naaggaaagg cggagcttat 180
tcaaagtcta gaggagtg aggagttang gctggatttc a 221

```

&lt;210&gt; 391

&lt;211&gt; 325

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1) ... (325)  
 <223> n = A,T,C or G

<400> 391  
 tggagcaggt cccgaggcct ccttagagcc tggggccgac tctgtgncga tgcangcttt 60  
 ctctcgcccc cagcctggag ctgtccctgg catctacca CActcagncg aggcgagcag 120  
 tagccragggc actgctgccc acagccagtc cnnatcccat catgtacccc ggtgngctct 180  
 naanttgat ntccanagcc ctaccatcn tagttctgct ctcccaccgg ntaccagccc 240  
 cactgccag gaa:cttaca gcagtagcc tgtcccagcg tctctacctc ccagtacgat 300  
 gggacctccg gctactacta tgacc 325

<210> 392  
 <211> 277  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1) ... (277)  
 <223> n = A,T,C or G

<400> 392  
 atallgctta actccttctt ctatatcttt caacattttc atggngaaug gttccacatcc 60  
 agtctcactt nggcncagga ctctactctg agtctcttcc cgggctggn ccagtngnaa 120  
 antaccanga accgncatgn cttaanaacc ncttggtttn tgggttuntc aatgar:gc 180  
 tgcagtgcac caccctgtcc actacgtgat gctgtaggat caaagcttca cagtggcgcg 240  
 ctgaggatcc agggcccgct cctgtgttgc tggggaa 277

<210> 393  
 <211> 566  
 <212> DNA  
 <213> Homo sapiens

<400> 393  
 actagtcacg tgtggtggaa ttccggggcc cgtcgacgga caggtcagct gtctggctca 60  
 gtgatctaca ttctgaagtt gtctgaaat gtcttcatga ttcaattcag cctaaacggt 120  
 ttgcccggaa cactgcagag acaatgctgt gagtttccaa ccttagccca tctgcgggca 180  
 gagaaggtct agtttgtcca tcagcattat catgatctca ggaactggtta cttagttaag 240  
 gagggtctc ggaatctgt cctttttaga gaccccttac ttataatgaa gtatttggga 300  
 ggggtgttt caaaagtga aatgtcctgt attccgatga tcatcctgta aacattttat 360  
 catttattaa tcatccctgc ctgtgtctat tattatattc atctctctac gctggaact 420  
 ttctgectca atgtttactg tgcctttgtt ttactagtt tgtgtgttg aaaaaaaa 480  
 cattctctgc ctgagtttca atttttgtcc aaagttattt taatctatac aattaaagc 540  
 ttttgcctat ccaaaaaaaa aaaaa 566

<210> 394  
 <211> 384  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature

<222> (1)...(384)

<223> n = A,T,C or G

<400> 394

```

gaacatacat gtccccggcgc ctgagctgca gtctgacatc atcgccatca cgggctctgc 60
tgcaaattns gaccggggcna aggcctggat gctggagcgt gtggaaggagc tacaggccna 120
gcaggaggag cgggctttta ggagttttta gctgagtgto actgtagacc ccaatataca 180
tcucaagatt atcggggagaa aggggggcagt aattacccaa atccggttgg agcatgacgt 240
gaacatccag tttcctgata aggacgatgg gaaccagccc caggaccana ttaucataca 300
agggtaacga aagaacacag aagctgcacg ggatgctata ctgagaattg tgggtgaact 360
tgagcagatg gtttttgagg acgt

```

384

<210> 395

<211> 399

<212> DNA

<213> Homo sapiens

<400> 395

```

ggcaaaactg tgtgacctca ataagacctc gcagatccaa ggtcaagtat cgggaagtgc 60
tctgaccttg gactccaaga cctacatcaa cagcctggct atattagatg atgagccagt 120
tatcagaggt ttcatcattg cgggaattgt ggagtcbaag gaaatcatgg cctctgaagt 180
attcagctct ttcaggtacc ctgagttctc tatagagttg cctaacacag gcgaattgg 240
ccagctactt gtctgcaatc gtatcttcaa gaatacctg gcaatccctt tgactgacgt 300
caagttctct tgggaagcc tgggcattct ctcactacag acctctgacc atgggacggt 360
gcagcctggt gagaccatcc aatcccaaat aaatgcaac

```

399

<210> 396

<211> 403

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(403)

<223> n = A,T,C or G

<400> 396

```

tggagttntc agtgraaaca agccataaag cttcagtagc aaattactgt ctacacagaa 60
gacattttca actttgtctc cagctgctga taaaacaaat catgtgctta gcttgactcc 120
agacaaggac aacctgttcc ttcataaotc tctagagaaa aaaaggagtt gttagtayat 180
actaaaaaaa gtggatgaat aatctggata tttttcctaa aaagattcct tgaaacacat 240
taggaaaatg gagggccttc tgatcagaat gctagaatta gtccattgtg ctggaagcag 300
gttttagggg yggagtggg gataaagaa gyaazaaag aagagtgaag aaacctattt 360
atcaaagcag gtgctatcac tcaatgttag gccctgctct ttt

```

403

<210> 397

<211> 400

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(100)

<223> n = A,T,C or G

&lt;400&gt; 397

actagtnrag tgtgggtggas ttgcggggcg cgtgcgccta naanccatct ctctagcaaa 60  
tccatccccg ctctgggttg gtnaccgaat gactgacaaa 100

&lt;210&gt; 398

&lt;211&gt; 278

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (278)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 398

gcggcgcggt cgaragcagt tccgcacagc ctgcgccttg ggtggggatg tgcgcacgc 60  
ccacctggac atctggaggt cagcggcctg gatgaagag cggacttcac ctggggcgat 120  
tcactactgt gctcgcaca gtggggagag ctggaccgac agcgaggttg actcactatg 180  
ctcggggcag cccatccacc tgtggcaglt cctcaggag ttgctactca agccaccagc 240  
ctatggccgc ttcattangt ggtccancaa ggagaagg 278

&lt;210&gt; 399

&lt;211&gt; 298

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)... (298)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 399

acgggggttg aggaagcgnc cctgggagtc anaggatggg tctcgcatt gaccncccn 60  
ggggtgccng catggagcgc atgggcgcgg gctggggcca cggcatggat cagctgggt 120  
ccgagatcga ggcgatgggc ctggtcatgg accgcattgg ctccctggag cgratgggc 180  
ccggcattga ggcgatgggc ccgctgggac tgcaccacat ggctccanc attganccga 240  
tgggcagac catggagcgc attggctctg gcgtggagcn catgggtgac ggcattggg 298

&lt;210&gt; 400

&lt;211&gt; 548

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 400

acatcaacta cttcctcatt ttaaggatag gcagttcctt tcctcccttt ttctgctt 60  
gtacatgtac atgtatgaaa tttccttctc ttaccgaact ctctccacac atcacagggt 120  
caaagaacca cagccttaga agggtaagag ggcacccctat gaaatgaat ggtgatttct 180  
tgagtctctt tttccacgt ttaaggggcr atggcaggac ttgaggttgc gagttaagac 240  
tgcagagggc tagagaatta tttcatacag gctttgaggc caccatgtu atttatccng 300  
tatccctctt caccatccc ttgtctactc tgatgccccc aagatgcaac tgggcagcta 360  
gttggcccca taattctggg cctttgttgt ttgttttaac tacttgggca tcccaggagc 420  
ctttccagtg atctctaac atgggccccc ctctgggat caagccctc ccaggccctg 480  
tcccagccc ctcctgccc agcccacccg ottgccttgg tgcctagccc tccatttggg 540  
agcaggtt 548

<210> 401  
 <211> 355  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(355)  
 <223> n = A,T,C or G

<400> 401  
 actgtttcca tgttatgttt ctacacattg ctacctcagt gctcctggaa acttagcttt 60  
 tgaatgtctcc aagtagtcca ctttcattta actctttgaa actgtatcat ctttgccaaag 120  
 taagagtggc ggcctatttc agctgctttg acaaaatgac tggctcctga ctttaacgttc 180  
 tataaatgaa tgtgctgaag caaagtgcgc atggtggcgg cgaagaagan aagatgtgt 240  
 tttgttttgg actctctgtg gtctcttcca atgctgmggg tttccaaaca ggggaagggt 300  
 cctttttgca ttgccaagtg ccataacctat gaggactact ctaccatggg tctgc 355

<210> 402  
 <211> 407  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(407)  
 <223> n = A,T,C or G

<400> 402  
 atggggcaag ctggataaag aaccaagacc cactggagta tgcgtctctc aagaaaccca 60  
 tctcactgc ggtggcctac ataggctcaa actaaaggaa tggagaaaaa tctttcaagc 120  
 aatggaaaa cagaaaaaag caggtgttgc actcctactt tctgacaaaa cagactatgc 180  
 gaataaagat aaaaaagaga aggacattac aaaggtggtc ctgacctttg ataaatctca 240  
 ttgcttgata ccaacccggg ctgttttaant tgcacaaerc aaaggatca tttgtgag 300  
 ttgtgaggt tctccctgc agagagtccc tcatctccca aaatttgggt gagatgtaag 360  
 gntgattttg ctgacaaetc ctttcttgaa gtttactcca ctccaa 407

<210> 403  
 <211> 303  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(303)  
 <223> n = A,T,C or G

<400> 403  
 cagtatttat agcnaactg aaaagctagt agcaggcag tctcaaatcc aggcaccana 60  
 tcttaagcaa gaggcatggc atggtgaaa tgcaaaagga gactctggcc aatctacaaa 120  
 tagagaacaa gacctactca gtcatgaaca aaaggcgaga caccacatg gatctcatgg 180  
 gggattggat attgttaata tagagcagga agatgacagt gctcgtcatt tggcacaaca 240  
 tcttaacaaac gaccgaaacc catttatlca ataaacctcc attcggtaac catgttgaaa 300  
 gga 303

<210> 404  
 <211> 225  
 <212> DNA  
 <213> Homo sapiens

<400> 404  
 aagtgttaact tttaaaattt tagtggattt tgaataattct tagagggaag taaaggaaaa 60  
 attgttaattg caatcattta cttttacatg gtgaaggctt tctcttgatc ctacaaacag 120  
 acattttcca ctctgtgtttc catagtgttt aagtgtatca gatgtgttg gcattgtgaat 180  
 ctccaagtgc ctgtgttaata aataaagtar ctttatttva ttrct 225

<210> 405  
 <211> 334  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(334)  
 <223> n = A,T,C or G

<400> 405  
 gagctgttct actgtgagtt ctactaggaa atcatcaaat ctgagggttg tctggaggac 60  
 ttcaatacac ctccccccat agtgaatcag ctccaggagg gtccagtccc tctcttaact 120  
 tcatccccat cccatgccaa aggaagaccr tccctccttg gctcacagcc ctctctagge 180  
 tccccagtgc ctccaggaca gactgggtta tgttttcagc tccatccttg ctgtgagtgt 240  
 ctggtgcggt tgtgactcca gcttctgctc agtgcctcat ggcagtgct cagcccatgt 300  
 caetctccac tctctcann tggaaccac cact 334

<210> 406  
 <211> 216  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(216)  
 <223> n = A,T,C or G

<400> 406  
 ttccatacct aatgagggag ttganatnac atnaaccrag gaattgcatt gatctcaang 60  
 gaaacannca cccaataaac tcggagtggt agactgacaa ctgtgagaca tgcatttgc 120  
 acnaaaraca aattttatgt tgcacccttg tttctacacc tgtgggttat gacaaagaca 180  
 actgocaaag aatnttcaag aaggaggact gacant 216

<210> 407  
 <211> 413  
 <212> DNA  
 <213> Homo sapiens

<400> 407  
 gctgacttgc tagtatcatt tgcattcatt gaagacacag aacttcatgc cttyactcat 60  
 gtaaatgraa taggattaaa aaataaattt gatattcac. ggaacagac aaaaaatatt 120  
 gtacaacatt gcacccagtg tcagattcta vacctggcca ctgaggagc agagattaat 180  
 cccagaggtc tatgtctta tatgtttatgg caaatggatg tcatgacgt accttcattt 240

```

ggagaaattgt cttttgtcca tgtgacagtt gatacttatt cacatttcac atggggcaacc 300
tgccagacag gagaaagtct tcccatgtta aaagacattt attatcttgt tctcctgtca 360
tgggagttcc agaaaaagtt aaaacagaca atggggccagg ttctgtagta aag          413

```

```

<210> 408
<211> 183
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1) ... (183)
<223> n = A,T,C or G

```

```

<400> 408
ggagctngcc ctraattccct ccatntctat gttancatat ttaatgtctt ctymnattaa 60
tnccttaacta gttactcctt aaagggcten ntatctctta actagtcctt ccathgtgag 120
cattatcctt ccagtattcn ccttcntttt catttactcc ttcctggcta cccatgtact 180
ntt                                     183

```

```

<210> 409
<211> 250
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1) ... (250)
<223> n = A,T,C or G

```

```

<400> 409
cccacgcatg ataagctctt tatttctgta agtctgtcta ggaatcctc aaatctgacg 60
gtgggtttgg ggacctgaac aaacctcctg taattaatca gctttcagtt tctcccccta 120
gtcctctctt caacaacata ggaggatcct cccctctctt ctgctcacgg ccttatctag 180
gcttcccaat gcccccagga cagcgtgggc tatgtttacg gcgctcctt gctggggggg 240
ggcmtatgc                                     250

```

```

<210> 410
<211> 306
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1) ... (306)
<223> n = A,T,C or G

```

```

<400> 410
ggctgggttg caagantgaa atgaatgatt ctacagctag gacttaacct tgaatggaa 60
agtcttgcaa tccattttgc aggatccgtc tgtgacatg cctctgtaga gtagcagcatt 120
cccggggacc ttggaaacag ttggcctgtt aaggtgcttg cteccccaga cacatccctn 180
aaggtgttgt aatggcgaaa accgcttccct tctttattgc ccttctttat ttatgtgaac 240
nactggttgg ctttttttgn atcttcttta aactggaaag ttcaattgng aaaatgaata 300
tcntgc                                     306

```

<210> 411  
 <211> 261  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(261)  
 <223> n = A,T,C or G

<400> 411  
 agagatattt cttaggtnaa agttcataga gttcccatga actatatgar tggcaccaca 60  
 ggaatttttt tatttaagga ttctgagatt ttgttgagc aggattagat aaggctgttc 120  
 tttaaatgtc tgaatggaa cagatttcaa aaaaaaaccc caccatctag ggtgggaaac 180  
 aggaaggaaa gatgtgata ggcctgatgg ccaaaaacca atttaccat cagttcnagc 240  
 ctctctcaa ggngaggcaa a 261

<210> 412  
 <211> 241  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(241)  
 <223> n = A,T,C or G

<400> 412  
 gtccaatgtt acctgacatt tctacacac cccatccac gatgtattcg ttgcccagtg 60  
 ggaacatacc agcctgaatt tggaaaaaat aattgtgttt ctgcccagg aactactarg 120  
 actgaatttg atggctccac aacataaac cagtgtaaa acagaagatg tggaggggag 180  
 ctgggagatt tcatctggta cattgaattc ccaactaac cangcaatta cccagccaac 240  
 a 241

<210> 413  
 <211> 231  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(231)  
 <223> n = A,T,C or G

<400> 413  
 aactcttaca atccaagtga ctcatctgtg tgattgaatc cttccactg tctcatctcc 60  
 ctcatccaag tctctagtac ctctcttttg ttgtgaaggc taatcaact gaacacaaa 120  
 aagtttactc tctcatttg gaacctaaa actctcttct tctgggtct gagggtcc 180  
 agaatccttg aatcattct cagatcattg gggacaccan atcaggaacc t 231

<210> 414  
 <211> 234  
 <212> DNA  
 <213> Homo sapiens



&lt;400&gt; 414

actgtccatg aagcaactgag cagaagctgg aggcacacac caccagacac lccacgcaag 60  
 gatggagctg aacacataac ccaactctgc ctggaggcag tgggaagcct agagaaggct 120  
 gtgagccaa gagggaaggt cttcttttg catgggatyg ggatgaagta aggaagagga 180  
 ctggaccccc tggagctga ttcactatyg ggggaggtgt attgaagtc tca 234

&lt;210&gt; 415

&lt;211&gt; 217

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}... (217)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 415

gcataggatt aagactgagt atcttttcta cttcttttta actttctaag gggcacttct 60  
 caaaacacag accaggtagc aatctctcac tgcctaaagg ntctcacac cactttctca 120  
 cacttagcaa tagtagaatt cagtcctact tctgaggcca gaagaatggt tcaaaaat 180  
 antggaattat aaaaataac aattaagaaa aataatc 217

&lt;210&gt; 416

&lt;211&gt; 213

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}... (213)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 416

atgcataatc aaaggaact gctctgcttt tagaagacat ctggactgct ctctgcatga 60  
 ggcacagcag taagctctt tgaattccag aatcaagaac tctctcttc agactattac 120  
 cgaatgcaag gtggttaatt gaaggccact aattgatgt caaatagaag gatattgact 180  
 atattggac agatggagtc tctactacaa aag 213

&lt;210&gt; 417

&lt;211&gt; 303

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}... (303)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 417

lactctttag gcccatcagg gaagttcaca ctggagagaa gtcatacata tctactgtat 60  
 gtgggaaggg ctttactctg agttcaatc tccaagccca tcagagagtc cactctgag 120  
 agaagccata caaatgcant gagtgtggga agaggttcag gagggattcc cattatcag 180  
 ttcattctagt ggtcacaca ggaaggaac cctataaatg tgagatatgt ggggaagggt 240  
 tcaatcaag ttcgtatctt caatcctc aggaaggncc cagtatanen aaacctttta 300  
 agt 303

<210> 418  
 <211> 328  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(328)  
 <223> n = A,T,C or G

<400> 418  
 ttctttggcgg tggatgggga gggatgggac angagtctca ctctgttgc caggctggag 60  
 tgcacaggca tgatctggc tcactccaa cctgacctcc catgtccaa cgtattctgt 120  
 ggcctcggct tccctgtag tagaattaca ggcacatgcc accacacca gctagttttt 180  
 gtatttttag tagagacagg gtttcacct gttggcagg ctggcttcaa actcctnacc 240  
 tcagnggtca ggcctggctc aaactcctga cctcaagtga tctgccacc tggcctccc 300  
 aaagtgcctn gtttaraggg cgtgagcc 328

<210> 419  
 <211> 389  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(389)  
 <223> n = A,T,C or G

<400> 419  
 cctcctcaag acggcctgtg gtccgcctcc cggcaaccaa gaagcctgca glgcatatg 60  
 acccctgagc catggactgg agcctgaaag gcagcgtaca ccttgcctct gatcttgctg 120  
 ctctgttctc ctctgtggct ccattccatg caccgttgtt gcactgaggc ttgtgcagge 180  
 cgagcaaggc caagctggct caaagagcaa ccagtcaact ctgccacggt gtgccaggca 240  
 ccggttctcc agccaccaa ctcactcgtt cccgcgaatg gcacatcagt tctctatccc 300  
 taaggttagg accaaagggc atctgctttt ctgaagtcct ctgctctatc agccatracg 360  
 tggcagccac tcnngctgtg tgaacggg 389

<210> 420  
 <211> 408  
 <212> DNA  
 <213> Homo sapiens

<400> 420  
 gtctctccta actcctgcca gaaacagctc tctcaacat gagagctgca cccctcctcc 60  
 tggccagggc agcaagcctt agccttggct tcttgtttct gcttttttc tggctagacc 120  
 gaagtgtact agccaaggag ttgaagtctg tgactttggt gtttcggcat ggagaccgaa 180  
 gtcccattga cactttccc actgaccca taaggaatc ctcatggcca caaggatttg 240  
 gccaactcac ccagctgggc atygagcagc attatgaact tggagagtat ataagaaaga 300  
 gatatagaaa atttttgat gagtctata aacatgaaca ggtttatatt cgaagcacag 360  
 acgttgaccg gactttgatg aagtgcctatg acaaccctgg caagcccg 408

<210> 421  
 <211> 352  
 <212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1) ... (352)

<223> n = A,T,C or G

<400> 421

```
gctcaaaaat ctttttactg atnggcctgg ctacacaatc attgactatt acggaggcca 60
ggagaggaatg aggcctggcc tgggagccct gtgcctacta naagcacatt agattatcca 120
ttcactgaca ggaacaggtct ttttgggttc cttcttctcc accacnatac acttgcagtc 180
ctccttcttg aagattcttt ggcagttgtc tttgtcataa cccacaggtg tggaaacaag 240
gggtcaacat gaaatttctg tttcgtagca agtgcattgc tcaacaagtg gcaagtctgc 300
cactccagat ttattgggtg tttgttctct ttgagatcca tgcatttctt gg 352
```

<210> 422

<211> 337

<212> DNA

<213> Homo sapiens

<400> 422

```
atgccaccat gctggcaatg cagcgggagg tccaaggcct gcatatccag cccaagctgg 60
cgaatgatcga cggcaaccgt tgcocgaagt tgcgcatgca agccgaagcg gtggtrcaagg 120
gcgatagcaa ggtgcggggc atngcggcgg cgtcaatcct ggcgaaggtc agccgtgctc 180
gtgaaatggc agctgtcgaa ttgatctacc cgggttatgg catcgggagg cctaagggct 240
atccgacacc ggtgcacctg gaagccttgc agcggctggg gccgagggcg attcaccgac 300
gtttcttccg ccggtacggc tggctatga aaattat 337
```

<210> 423

<211> 310

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1) ... (310)

<223> n = A,T,C or G

<400> 423

```
gctcaaaaat ctttttactg atatggcatg gctacacaat cattgactat tagaggccag 60
aggagaatga ggcctggcct gggagccctg tgcctactan aagcncatta gattatccat 120
tcaactgacag aacaggtctt ttttgggttc cttcttctcc ccargatata cttgcagttc 180
tccttcttga agattctttg gcaattgtct ttgtcataac ccacaggtgt anaaacaaag 240
gtgcaacatg aaatttctgt ttcgtagcaa gtcgatgtct cacagttgtc aagtctgccc 300
tcaggattta 310
```

<210> 424

<211> 370

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1) ... (370)

<223> n = A,T,C or G

&lt;400&gt; 424

```

gctcaaaaat cttcttactg ataggcatgg ctacacatc attgactatt agaggucaga 60
ggagaatgag guctggcctg ggagccclgt gctactaga agcacattag attatccatt 120
cactgacaga acaggtcttt ttgggtctt ccttctccac cactatatac ttgcagtcct 180
ccttctttaa gattctttg cagttgtctt tgtcataacc cccagggtga gaacatcct 240
ggttgaatct cctggaaact cctcattagg tatgaatat catgatgcat tgcataaagt 300
cccgaaaggtt gcaagatca caacgctgct cagganaaca ttcattgtga taagcaggac 360
tcctgtagcg

```

370

&lt;210&gt; 425

&lt;211&gt; 216

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(216)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 425

```

aattgctatn ntttattttt ccaactcaaa caattaccaa aaaaaaanaa tnttaaatga 60
taacaaunaa acatcaaggc aaaaaaanaa ggaatggntg actntgcta aatngggcca 120
anattatcca ttatnttaa ggttgacttc aggtacagc acacagacaa acatgcccag 180
gaggntctca ggaacgctcg atgtntcttg aggagg

```

216

&lt;210&gt; 426

&lt;211&gt; 596

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 426

```

cttcagtgga ggataacccct gttgcrcagg gccgagggtc tccattaggc tctgattgat 60
tggcagtcag tgatggaagg gtgttctaat catccagact gcccaaggc tctctggcca 120
gctctctgtt tngctgagtt ggcagtagga cctaatctgt taattaagag tagatngtga 180
gctgtccctg tattttgatt aacctaatg ccttccagc ccgactcgga ttcagctgga 240
gacatcacgg caacttttaa tgaatgatt tgaagggtcc ttaagaggca ctcccggtta 300
ttaggcagtt catctgcact gataactct tggcagctga gctgttcgga gctgtggccc 360
aaacgcacac ttggcctttg gttttgagat caaactctta atcttctagt catgcttgag 420
ggfagatggc cttttcagct ttaacccaat ttgactgac ttggaagtgt agcaggagga 480
atacartcat atactcgtg gcttagaggc cacagcagat gtcattgggc taetgcttga 540
gtcccgctgg tccatccca ggaccttcca tgggcagta cctgggagcc cgtgct 596

```

&lt;210&gt; 427

&lt;211&gt; 107

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(107)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 427

```

gaagaattca agttagggtt attcaaggc ctlaengaga atcctanacc caggmuccag 60

```

cccgaggaga gccttanaga gctccctgtt gactgcccgg ctcaagg

107

<210> 428

<211> 38

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(38)

<223> n = A,T,C or G

<400> 428

gaatttcma anaaagactt tattcaatat ttacatt

38

<210> 429

<211> 544

<212> DNA

<213> Homo sapiens

<400> 429

ctttgctgga cggaaataas gtggaagcaa gcattgacct ctgatgaggg cgcctgcattt 60  
attgaagagc ggcctgcagcc ctgcgggttca gattaaaatc cgagaatttgt atagacgccg 120  
atatccacga actcttgaag gctttcttga ttatccaca atccaaatcat cggttttcag 180  
tttggatggg ggcctcatcc ctgtagaacc tgccttgccc gtggctggaa tccactcgtt 240  
gccttccact tcagttacac ctccctcacc atctctctct gttggctctg tgcctctca 300  
agatactaag cccacatttg agatgcagca gcaatctccc ccaattctct ctgtccatcc 360  
tgatgtgcag tcaaaaaatc tgccttttla tgaatgcttt gaggttctca tcaagccac 420  
gagtttagtt caaagcagta ttacagcatt tcaagagaa ttttttatt ttgctttgac 480  
acctcaacaa gttcagagaga tatgcatac cggggatttt ttgccaggty gtaggagaga 540  
ttat

544

<210> 430

<211> 507

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(507)

<223> n = A,T,C or G

<400> 430

cttctcncaa tggggctccc aaacttggt gtgcagtga aactccgggg gaattttgaa 60  
gaacactgac acccatcttc caccrcgara ctctgattta attgggctgc agtgagaara 120  
gagcatcaat ttaaaaagct gcccaagaatg ttntcctggg cagcgttgtg atcttctgcn 180  
ccttngtgac tttatgcaat gcataatgct atttcatacc taatgagggg gtccaggag 240  
attcaaccag gatgtttcta cncctgtggg ttatgacaaa gacaartgc aagcaatntt 300  
caagaaggag gactgcaagt atatcgtggt gygagagaag gacccaaaa agacctgtt 360  
tgtcagtga tggatctct aatgtgcttc tagtaggcac agggctcccc ggcaggcct 420  
cattctctct tggcctctaa tagtcaatga ttgtgtagcc atguctatca gtaaaaagat 480  
ttttgagcaa aaaaaa aaaaaa

507

<210> 431

<211> 392

<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(392)  
<223> n = A,T,C or G

<400> 431  
gaaattcag catggataaa aacaaatgaa gtacaaata ttccagattt aatagcgat 60  
aaacaagaaa gcacttatca gagggaactta caaatggaa tacaactctan aaccatcato 120  
taccatgget aatgtgaga ttagcacagc tgtattattt gtacattgca aacacctaga 180  
aagagatggg aaacaaatc ccaggagttt tgtgtgtggg gtccctgggtt tccaaacaga 240  
catcattcca gcattctgag attagggnga ttgggataca ttctggagtt ggaatgttca 300  
acaaagtgga tgttgttagg taaatgtac aacttctgga ttatgcraga cattgaagg 360  
gcaatgaatc tggcttttac tctgtgtt ct 392

<210> 432  
<211> 387  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(387)  
<223> n = A,T,C or G

<400> 432  
ggatccnta cabaetcaa tatagctgta gtacatgttt tcattggngt agattaccac 60  
aatgcaagg caacatgtgt agatctcttg ttttattctt ttgttatata tactgtattg 120  
ngtagtccaa gctctcggna gtccagccac tnggaacat gctcccttta gatlaaacctc 180  
gtggaacnctn ttgttgnatt gctggaacty tagngccctg ttttttctt ctgtctgnga 240  
attctgttcg ttctggggca ttctcttngg atgcagaagg ccaccacaca gatgacagca 300  
ctctgaattt ntccaatcac agctgcgatt aagacatact gaatcgtac aggaccggga 360  
acaacgtata gaacactgga gtccctt 387

<210> 433  
<211> 281  
<212> DNA  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<222> (1)...(281)  
<223> n = A,T,C or G

<400> 433  
ttcaactagc anagaaact gcttragggt ggtanattg aaaggcttcc acgagttat 60  
ctgattaaag aacactaaga gaggyncaag gctagaagcc gaaggatgtc tacactatag 120  
caggcnctat ttggttggc tggaggagcl gtggaaaca tggagagatt ggcgtggag 180  
atcgccgtgg ctattcctn ttgntattac accagnaggy ntctctgtnt gccactggt 240  
tnnaaaacgy ntatacaata atgatagaat aggacacaca t 281

<210> 434  
<211> 484

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 434

```

tttctaaata agcattttagt gctcagttcc tactgggtac tctttctctc ccttctctctg 60
aatttaattc ttccaacttg caabttgaaa ggattacaca tttcactctg atgtatatctg 120
tgtctgcaaaa aaaaaaagt gtctttgttt aaaaattact ggtctgtgaa tccatcttgc 180
tttttcccca ttggaactag tcatbaacac atctctgaac tggtagaaaa acatctgaag 240
agctagtcctc tcagcatctg acaggtgaat tggcgggttc tcagaacctt ttcacccaga 300
cagctctgtt ctatctgtt taataaatta gtttgggttc tctcatgca taaccaaacn 360
tgcctccatc tctcacata aagttctgtg cttagagttt agtcagcacc cccacccacc 420
tttatttttc tatggtttt tgcacacata tgagtgtttt gaaataaag taccatgtc 480
tcta

```

484

&lt;210&gt; 435

&lt;211&gt; 424

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 435

```

ggcgcgtctc ggcaggtcca cttcttgcct tccaggtcct ccttcaggga agcccatctg 60
gggtagcttt caatategca ggttcttact cctctgcttc tataagctca aacccacca 120
cgatcgggca agtaaacccc ctccctcgcc gacttcggaa ctggcgagag ttcagcgccg 180
atggggtctt ggggaggggg caagatagct ggggaggagc ggcattggtc ggggtgaccc 240
cttggagaga ggaataaggc cactagaggg gctgcccagg cactaacgg agatggccct 300
ggtagagacc tttaggggtc tggaaacctt ggaacctcca tgccttaact cccacatct 360
gctatcagaa acttaaaact ggggatttct tctgttttct actgcaata aattcagagc 420
aaac

```

424

&lt;210&gt; 436

&lt;211&gt; 667

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (667)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 436

```

accttgggaa nactctcaca atataaaggg tcttagactt tactccaaat tccaaaagg 60
tcttggcat gtaatcttga aagttttccc aaggtagcta taataacctt ataagggtgc 120
aguctcttct ggaattcttc tgatttcaaa gtctcactct caagttcttg aaaaagagg 180
cagttcttga aaggtcaggt tagcaactga tcttcagaaa gaggaactgt gtgcacccgg 240
atgggtctgc agagtaggct aggtttccag atgtgacac ctctcggggg aaacagggtc 300
gccaggtttg tctagcact catcaaagtc cagtcacagt ctgtgcttcg aatatcaacc 360
tgttcatgtt tataggactc attcaagaat tttctatctc tctttcttat atactctcra 420
agttctaat gctgtctcat gcccgctggg gtgagttggc caaatcttg tggccatgag 480
gattcttlla tgggtcaggt gggaaagggt tcaatggguc ttcgggtctc atgccgaac 540
accaaagtc caaacttcaa ctcttgggt agtaccttc ggtctagcaa gaaaaaagg 600
agaaacaaga agcaaaggct aaggcttgc tgcctggcag gaggaggggt gcagctctca 660
tgttag

```

667

&lt;210&gt; 437

&lt;211&gt; 693

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 437

```

ctacgtotca accctcattt ttaggtzagg aatcttaagl ccaagatata taagtgaetc 60
acacagccag gtaaggaaag ctggattggc acactaggac tctaccatac cgggttttqt 120
taagotcag gttaggaggc tgaataagct ggaaggaaet ccagacagct ttccagatc 180
ataaagata attcttager catgtctctc tccagagcag acctgaatg ccagcaccgc 240
aggtaectct ctattttcac cctctttgct tctactctct ggcagtcaga cctgtgggag 300
gccctgggag aaagcagctc tctggatglt tgaacagatc atggactatt ctctgtggac 360
cattttcca ggtcacctca ggtgtcacta ttggggggac agccagcacc tttagcttct 420
atttgagttt ctgtctgtct tcagtagagg aaacttttgc tcttcacact tccatclga 480
acacctaaet gctgttgctc ctgaggtggt gaaagacaga tatagagctt accgtattta 540
tctattttct aggaactgag gctgtgggg taccttggcg tgcnaaaaca gatcctgttt 600
taaggacatg ttgcttcaga gatgtctgta actatctggg ggtctctgtg gctctttacc 660
ctgcacatg tgtctctctg gctgaaatg acc

```

693

&lt;210&gt; 438

&lt;211&gt; 360

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 438

```

ctgottatca caatgaatgt tctctgggc agcgttgtaa tctttgccac ctctgtgact 60
ttatgcacag catcatgcta ttcatacct catgaggag ttccaggaga ttcaaccagg 120
atgtttctac acctgtgggt tatgcnaag acaactgcc aagatcttc aagaaggagg 180
actgraagta tatctggtg agagaaggc ccaaaaaag acctgttctg tcagtgaatg 240
gataatctaa tgtgtttcta gtaggacag ggtcccag ccaggcctca ttctctctg 300
gctctataa gtcaataatt gtgtagccat gcttatcagt naaaagattt ttgagcaaac 360

```

&lt;210&gt; 439

&lt;211&gt; 431

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(431)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 439

```

gttccctnta actcctgcc aaaaagctc tctcaacat gagagctgca cccctctcc 60
tggccagggc agcaagcctt agccttggct tcttgtttct gcttttttct tggctagacc 120
gaagtgtact agccaaggag tgaagtctg tgactttggt atttcggcat ggagacgaa 180
gtcccattga cacttttccc actgacccca taagggaatc ctcatggcca caaggatttg 240
gccaactcac ccagctgggc atggagcagr attatgaact tggagagtar ataagaaaga 300
gatatagaan attcttgaat gactcctata aacatgaaca ggtttatatt cgaagcacng 360
acgttgaccg gactttgatg agtgctatga caaacctggc agccctcga cgcggccgag 420
aatttagtag t

```

431

&lt;210&gt; 440

&lt;211&gt; 523

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens



&lt;400&gt; 440

```

agagatcaag cttaggtcaa agttcataga gtteccatga actatctgac tggccacaca 60
ggatcttttg tatttaagga ttctgagatt ttgcttgggc aggattagat aaggctgttc 120
tttaaatgtc tgaatggaa cagatttcaa aaaaaaacc cacaatctag ggtgggaaca 180
aygaaggaa gatgtgaata ggctgatggg caaaaaacca atttaccat cagttccagc 240
cttctctcaa ggagaggcaa agaaaggaga taaggtggag acatctggaa agttctctcc 300
actggaaac tgcactatc tgttttata tttctgttaa atatctagag gttacagaac 360
taaaattca aactctcttg tgttcttgg tcttggaaac ttatcttcc ttttaagaa 420
acaaaatca aactcttca agagatttga tctatgtat acatatagca gctcttgaag 480
tatatatct atagcaata agtcatctga tggagacaag cta
523

```

&lt;210&gt; 441

&lt;211&gt; 430

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 441

```

gttcttctca actcctgcaa gaaacagctc tctcaacat gagagctgca cccctctcc 60
tggccagggc agcaagcctt agccttggct tcttgttct gtttttttc tgggtagacc 120
gaagtgtact agccaaggag ttgaagtttg tgaacttggg gtttggcat ggagacgaa 180
gtcccatgga cactttccc actgaaccca taagggaatc ctcatggca caaggatttg 240
gccaactcac ccagctgggc atgggacgac attatgaact tggagagtat atagaaaga 300
gatctagaaa attcttgaat gactcctata aacatgaaca ggtttatatt cgaagcacag 360
acgttgacgg gactttgatg agtgcctatga caaacctggc agccctcga cggggccggc 420
aatttctgag
430

```

&lt;210&gt; 442

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 442

```

ctaaggaatt agtagtgtc ccctcacttg ttggagtyt gctattctaa agattttga 60
tttcttggaa tgaacattat atttcaact tgggtgggga aagagttata ggaccacagt 120
cttcaattct gatacttgta attaatctt ttattgaact tgttttgacc attagctat 180
atgtttagaa atggtcattt tacggaaaaa ttgaaaaat tctgataata gtgcagaata 240
aatgaattaa tgttttactt attttatatt gaactgtcaa tgacaaataa aaattctttt 300
tgattatttt ttgttttcat ttaccagaat aaactaag aattaaagt ttgatcacag 360
tc
362

```

&lt;210&gt; 443

&lt;211&gt; 624

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1) ... (624)

&lt;223&gt; n = A, T, C or G

&lt;400&gt; 443

```

tttttttttt gcaacacaat atacatcaca gtgaatgtg caatccttgc aaattgcaag 60
ttgaagaaat taatatcaga ggaggggaga gaagagctac tcagtaggga ctgagcacta 120
aatgcttatt ttanngaaa tgtaaagagc agaaagcaat tcaagctacc ctgcttttg 180
tgcctgctag tactcgggtc ggtgtcagca gcacgtggaa ttgaacattg caatgtggag 240

```

```

cccaaacccac agaaaatggg gtgaatctgg ccaactttct attaacttgg ctccctgttt 300
tataaattat tgtgaataat atcacctact tcaaggggca gttatgaggg ttatatganc 360
taacgctac aaaaactta aacatagata acatagggtg aagtactatg tatctggtar 420
atggttaaca teottattat caaagtcacac gctaaatga atgtgtgtgc atatgcta 480
agtacagaga gagggcactt aaacccaacta agggcctgga ggggaaggltt cctggaaaga 540
ngatgcttgt gctgggtcca aatcttgggt tactalgaac ttggccaaat tatttaaa 600
ttgtccctat ctgctaaca gac 624

```

&lt;210&gt; 444

&lt;211&gt; 425

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(425)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 444

```

gcacatcatt nntcttgcct tctttgagaa taagaagatc agtaaatagt tcagaagctg 60
gaagctttgt ccaggcctgt gtgtgaaccc aatgttttgc ttagaatatg acaagtaag 120
ttcattgcta tagcataaca caaanatttg ataagtgggt gtacgcaaat ccttgaalgc 180
tgcttaattgt gagaggttgg taaaatcctt tgtgcacac tctaactccc tgaatgtttt 240
gctgtgctgg gacctgtgca tgcagacaa ggccaagctg gctgaagag caaccagcca 300
cctctgcaat ctgccacctc ctgctggcag gatttctttt tgcacctgt gaagagccaa 360
ggaggcacca gggcataagt gagtngactt atggctcgacg cggccgcgaa cttagtagta 420
gtaga 425

```

&lt;210&gt; 445

&lt;211&gt; 414

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(414)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 445

```

catgtttatg nttttggatt actttgggca cctagtgttt ctaaatcgta tatcattctt 60
ttctgttttt caaagcaga gatggccaga gtctcaacaa actgtatctt caagtctttg 120
tgaattctt tgcattgtgg agattattgg atgtagtctt ctttaactag catataaatc 180
tgggtgtttt cagataantg aacagcaaaa tgtgggtggaa ttacattttg gaacattgtg 240
aatgaaaaat tgtgtctcta gattatgtaa caaataacta ttccctaacc attgatcttt 300
ggatttttat aatcctatc acaaatgact aggtctctcc tcttgtatct tgaagcagtg 360
tgggtgctgg attgataaaa aaaaaaaag tcgagcgggc cgggaattta gtat 414

```

&lt;210&gt; 446

&lt;211&gt; 631

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(631)

<223> n = A,T,C or G

<400> 446

```

acaaattaga anaaagtguu agagaaracc acataacttg tccggacat tacaatggct 60
tctgcatgca tgggaagtgt gagcattcta tcaatatgca ggagccalcl tgcagggtgtg 120
atgctgggtta tactggaca cactgtgaaa aaaaggacta cagtgttcta tangltgttc 180
ccggtcctgt acgatttcag tagtctttaa tccgagctgt gattgggaca attcagattg 240
ctgtcatctg tctggtggtc ctctgcatca caagggccaa actttaggta atagcatttg 300
actgagattt gtaaaccttc caaccttcaa ggaatgccc cagaagcaac agaatccaca 360
gacagagcgc aatatcaggg cactacagtt cagacaalac aacagagcgc tccacagggc 420
taattcaag ggagcatgtt ccacagtgc tggactaccg agagcttggc utacacaata 480
cagtatctat gacaaagaa taagacaaga gatctacaca ttttgccttg catttctgtg 540
natctacacc aatgaaaaca tgtactacag ctatatttga ttatglatgg utatatttga 600
aatagtatac attgtcttga tgtttttct g 631

```

<210> 447

<211> 585

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> {1}... (585)

<223> n = A,T,C or G

<400> 447

```

ccttgggaaa antnlcaca tatnaaggt cgtagactcl actccaaatt ccaaaagggt 60
cctggccatg taatcctgaa agttttccc aagtagctat aaaatcctta taaagggtgca 120
gctctctctg gaattcctct gatttcaag tctcactcnc aagttcttga aaacgagggc 180
agtctctgan aggcagggtat agcaactgat ctccagaaag aggaactgtg tgcacuggga 240
tgggctgcca gactaggata ggaattccga tgcagacac ttctggggga aacagggtgt 300
ccaggcttgt catagacctc atcaaggtcc ggtcaacgtc tgtgcttcca atataaact 360
gttcctgttt ataggactca ttcaagaaat ttctatatct ctctcttata tactctccaa 420
gttcataatg ctgtccatg ccagctggg tgagtgggc aatccttgt ggccatgagg 480
attcctttat ggggtcagtg ggaaggtgt caatgggact tgggtctcca tgcrgaaaca 540
ccaaagtcac aaattcaac tcttggcta gacacttgc gtcta 585

```

<210> 448

<211> 93

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> {1}... (93)

<223> n = A,T,C or G

<400> 448

```

tactgttggg tctttctgan nccggaactg accttgcag ccttgcgaa gggucnccat 60
ggctccctag tgccttggag aggaaggggc tag 93

```

<210> 449

<211> 706

<212> DNA

<213> Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(706)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 449

```

ccaagttcat gctntgtgct ggacgctgga cagggggcaa aagcnnntgc tegtgggtca 60
ttctgancac cgaactgacc atgccagccc tgccgatggt cctccatggc tccctagtgc 120
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aacaggttga acctgggagg tgggagttgc aatgagctga gatcagggcn ctgmcacca 660
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```

&lt;210&gt; 450

&lt;211&gt; 493

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 450

```

gagacggagt gtcactctgt tgcccaggct ggagtgagc aagacactgt ctaagaaaa 60
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taccatcag aatracctgg agagutttac aaactcccat tgcgagggt cgagcggcc 480
gggaatttag tag

```

493

&lt;210&gt; 451

&lt;211&gt; 501

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(501)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 451

```

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ctcttcgcta ttacgccage tggcgaaagg gggatgtgt gcaaggcgat taagtgggt 120
aacgccaggg ttttcacagt cncgacgttg taaaacgac gcacgtgaa tgaattttag 180
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cgcnccagac actacagct actcaggagg ctgagacacg attgaacctg ggaggtgga 420
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```

tcttaaaaaa aaaaaaa a

501

<210> 452

<211> 51

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(51)

<223> n = A,T,C or G

<400> 452

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51

<210> 453

<211> 317

<212> DNA

<213> Homo sapiens

<220>

<221> misc\_feature

<222> (1)...(317)

<223> n = A,T,C or G

<400> 453

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ttcaccacac cagcctgttt ctatcctgtt caataaatta gtttgggttc tctaatgca 180  
taacaaaccc tgcaccaatc tgcacacata aagtctgtga cttgaagtt antcagacc 240  
cccaccacac tttatttttc tatgtgtttt ttgaacata tgaagtgttt gaaataagg 300  
taaccatgtc tttatta 317

<210> 454

<211> 231

<212> DNA

<213> Homo sapiens

<400> 454

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agaagaccac attcttctgc atccacagctt gcanacaaaa ttgtctctct aggtctccac 180  
ccttcttttt tcaagtgttc aaagctcctc acaatttcat gaacaacagc t 231

<210> 455

<211> 231

<212> DNA

<213> Homo sapiens

<400> 455

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gtttcaacgc attgatgact tctccaagga tcttctttg gcatcgacca cattcaggg 180  
caaagaattt ctcatagcac agctcaaat acagggtcc tttctctct a 231

<210> 456  
 <211> 231  
 <212> DNA  
 <213> Homo sapiens

<400> 456  
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 tgcactcaaa ttcttttatc aggaataact acctagccc cttttacaaa ggcattggaa 180  
 cctttttatt tggtagagct gctcgtcagt cctgactga cattgccag t 231

<210> 457  
 <211> 231  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> misc\_feature  
 <222> (1)...(231)  
 <223> N - A,T,C or G

<400> 457  
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 catttgattt tattagcaat ctctttcaga agaccttga gatcattag ctttgtatcc 180  
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<210> 458  
 <211> 231  
 <212> DNA  
 <213> Homo sapiens

<400> 458  
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 acacctaac cttgggtaac agcatttggg attatcattt ggyatgagta gaatttcaa 180  
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<210> 459  
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 <212> DNA  
 <213> Homo sapiens

<400> 459  
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 gccctgcact gttttccctc caccacagcc atctctgccc tctttggctc tgtgctttcc 180  
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<210> 460  
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 <212> DNA  
 <213> Homo sapiens

<400> 460

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ccnacctccc caccacgaca cggccagcct ggagcccaaca gaagggtcct cctgcagcca 180
gtggagcttg gtccagcctc caglcacccc ctaccaggct taaggataga a 231

```

&lt;210&gt; 461

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 461

```

cgagggttga gaagctctaa tgtgcaaggg agccgagaag caggcggcct agggagggtc 60
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gtgggggttca gtgaggagtg ggaatttggg ttagcagaac caagccgttg ggtgaataag 180
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```

&lt;210&gt; 462

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 462

```

aggtaccctc attgtagcca tgggaaaatt gatgttcagt ggggatcagt gaattaaatg 60
gggtcatgca agtataaaaa ttaaaaaaaa aagacttcat gcccaatctc atatgatgta 120
gaagaactgt tagagagacc caacagggtag tgggttagag atttcagag tcttaccttt 180
tctagaggag gtatttaatt ttttctcant cctccagtgt tgtatttgg a 231

```

&lt;210&gt; 463

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 463

```

tactccagcc tggtagacaga gggagaccct atcacggccc cccacccccc caaaaaaaan 60
actgagtaga caggtgtcct cttggcatgg taagtcttaa gtccctccc agatctgtga 120
catttgacag gtgtctttc ctctggacct cgggtgtccc atctgagtga gaagaagcag 180
tggggaggtg gatcttccag tcgaagcggc atagaagccc gtgtgaaaag c 231

```

&lt;210&gt; 464

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 464

```

gtactctaa agtttatcta agctgccttt tctgggtggg aaagtctaac cttagtgact 60
aaggacatca catatgaaga atgtttaaag tggaggtggc aacgtgaatt gcaaacaggg 120
cctgcttcag tgactgtatg cctgtagtc cagctacttg ggagtctgtg tgaggccagg 180
gggtccagcg caccagctag atgtctctga acttctaggc cccattttcc c 231

```

&lt;210&gt; 465

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 465

```

calgtttgttg tagctgtggt aatgctgggt gcatctcaga cagggttaac ttcagctcct 60
gtggcaaat agcaacaaat ttgacatca tatttatggt ttctgtatct ttgttgatga 120
aggatggcau aatttttgot tgggtt.cala atatactcag attagttcag ctucatcaga 180
taaactggag acatgcagga ctttagggta gtgttgtagc tctegtantg a 231

```

&lt;210&gt; 466

&lt;211&gt; 231

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 466

```

caggtaacct tttccattgg atactgtgct agcaagcatg ctctccgggg tttttttaat 60
ggccttcgaa cagaacttgc cacatcccc agtatantag ttctaacct ttgccagga 120
octgtgcaat caaatattgt gggaatttc ctagctggag aagtcacaaa gacttatggc 180
aatcatggag accagterca caagatgaca accagtgcgt gtgtcgggct g 231

```

&lt;210&gt; 467

&lt;211&gt; 311

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 467

```

gtacaccctg gcacagtcac atctgaactg gttcggcact catcttcat gagatggatg 60
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ctgcagcaga c 311

```

&lt;210&gt; 468

&lt;211&gt; 312

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 468

```

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```

&lt;210&gt; 469

&lt;211&gt; 2229

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 469

```

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&lt;210&gt; 470

&lt;211&gt; 2426

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 470

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2426

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&lt;210&gt; 471

&lt;211&gt; 812

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 471

```

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gagatcagat attacacag ctttgttttg agggtagaa atatgaastg atttgggtat 180
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812

```

&lt;210&gt; 472

&lt;211&gt; 515

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; misc\_feature

&lt;222&gt; {1}... (515)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 472

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ctctgatgta aaagagaaag ctcacctaac tcttaagcat gctaaagact aagaaagcag 360
agaggggcag ccataagtta aaagaagac aagctgaagc lacacacatg gctgatgctc 420

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Cattgaaat gtgactgaa attgaaat tctctcaata aagtttgagt tttctctgaa 480  
gaaaaaaa aaagaaaaa aaaaaaa aaaa 515